From:

Huff, Sheela

Sent:

Monday, September 13, 2004 5:58 AM

To: Subject: STIC-Biotech/ChemLib search request for 10042347

Please search and interference search SEQ ID No. 4 of the above application.

Thanks

Sheela Huff
Art Unit 1642
571-272-0834
Remsen 3A15
mailbox Remsen 3C18

STAFF USE ONLY
Searcher:

Searcher:
Searcher Phone: 2Date Searcher Picked up:
Date Completed:
Searcher Prep/Rev. Time:
Online Time:

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Type of Search
NA Sequence: #
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US-09-561-506-12

US-09-561-566-12

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US-09-561-267-12

US-09-561-77D-5

US-09-211-077D-7

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SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 4
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CURRENT APPLICATION NUMBER: US/09/315,689
CURRENT FILING DATE: 1999-05-20
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APPLICANT: O'Reilly, Michael
TITLE OF INVENTION: Therapeutic Antiangiogenic Endostatin Compositions
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LENGTH: 552
TYPE: DNA
ORGANISM: Homo sapiens
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APPLICANT: Sim, Kim Lee
TITLE OF INVENTION: Anglogenesis-Inhibiting Protein Binding Peptides and
TITLE OF INVENTION: Proteins and Methods of Use
FILE REFERENCE: 05213-0370
CURRENT APPLICATION NUMBER: US/09/206,059
CURRENT FILING DATE: 1998-12-04
NUMBER OF SEQ ID NOS: 80
SOFTWARE: Patentin Ver. 2.0
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Patent No. 6201104
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; ORGANISM: human
US-09-231-077D-9
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APPLICANT: Violand, B.N.
TITLE OF INVENTION: Method of producing mouse
TITLE OF INVENTION: endostatin
FILE REFERENCE: S03071-00-US
CURRENT APPLICATION NUMBER: US/09/231,077D
CURRENT FILING DATE: 1999-01-14
PRIOR APPLICATION NUMBER: 60/075,587
PRIOR FILING DATE: 1998-02-23
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Best Local Similarity
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
COMPUTER: IBM PS/2 Model 50Z or 55SX
OPERATING SYSTEM: MS-DOS (Version 5.0)
SOFTWARE: WordPerfect (Version 5.1)
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CORRESPONDENCE ADDRESS:
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VENTION: NOVEL COLLAGEN AND USES
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Pred. No. 1e-112;
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CURRENT FILING DATE: 1999-05-20
NUMBER OF SEQ ID NOS: 6
SOFTWARE: Patentin Ver: 2.0
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APPLICANT: O'Reilly, Michael
TITLE OF INVENTION: Therapeutic Antiangiogenic Endostatin Compositions
FILE REFERENCE: 05213-0229
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TYPE: DNA
ORGANISM: Homo sapiens
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                TGCGATCACGCCTACATCGTGCTCTGCATTGAGAACAGCTTCATGACTGCCTCC 546
                                                                                     GCCACGGGCCAGGCCTCCTCGCTGCTGGGGGGCAGGCTCCTGGGGCAGAGTGCCGCGAGC
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TGCCATCACGCCTACATCGTGCTCTGCATTGAGAACAGCTTCATGACTGCCTCC
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; TYPE: DNA
; ORGANISM: Rattus rattus
US-09-449-293-3
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APPLICANT: Uteza, Yves
APPLICANT: Menasche, Maurice
APPLICANT: Bossard, Carine
APPLICANT: Van Den Berghe, Loic
APPLICANT: Bonnel, Sebastien
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Best Local :
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SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 3
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APPLICANT: Neuner-Jehle, Martin
TITLE OF INVENTION: INTRAOCULAR TRANSPLANTATION OF ENCAPSULATED CELLS
FILE REFERENCE: 8076.202US01
CURRENT APPLICATION UNMEER: US/09/449,293
CURRENT FILING DATE: 1999-11-24
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TCTTTC
                                                      AAAGCTGCGAGCTGCACAACAGCTACATCGTCCTGTGCATTGAGAATAGCTTCATGACC
                                                                          AGTGCCGCGAGCTGCCATCACGCCTACATCGTGCTCTGCATTGAGAACAGCTTCATGACT 540
                                                                                                                                         CACGGCTCGGACCCCAGTGGGCGGAGGCTGATGGAGAGTTACTGTGAGACATGGCGAACT
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Prats, Herve
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; ORGANISM: Rattus rattus
US-09-775-325-3
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APPLICANT: Abitbol, Marc
APPLICANT: Uteza, Yves
APPLICANT: Menasche, Mau
APPLICANT: Mossard, Cari
APPLICANT: Mossard, Cari
APPLICANT: Wan Den Bergh
APPLICANT: Bonnel, Sebas
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// Patent No. 6500449
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Best Local Similarity
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APPLICANT: Neumer-Jehle, Martin
TITLE OF INVENTION: INTRACCULAR TRANSPLANTATION OF ENCAPSULATED CELLS
FILE REFERENCE: 8076.2020SD1
CURRENT APPLICATION NUMBER: US/09/775,325
CURRENT FILING DATE: 2001-02-01
PRIOR APPLICATION NUMBER: US 09/449,293
PRIOR PILING DATE: 1999-11-24
NUMBER OF SEQ ID NOS: 5
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                            AAAGCTGCGAGCTGCCACAACAGCTACATCGTCCTGTGCATTGAGAATAGCTTCATGACC 543
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Bossard, Carine
Van Den Berghe, Loic
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Prats, Herve
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Pred. No. 3.1e-76;
0; Mismatches 103;
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CURRENT FILING DATE: 2001-02-01
PRIOR APPLICATION NUMBER: US 09/449,293
PRIOR FILING DATE: 1999-11-24
NUMBER OF SEQ ID NOS: 5
SOFTWARE: PatentIn version 3.1
SEQ ID NO 3
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APPLICANT: Neuner-Jehle, Martin
TITLE OF INVENTION: INVEACCULAR TRANSPLANTATION OF ENCAPSULATED CELLS
FILE REFERENCE: 8076.202USD2
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                                                                                                AGTGCCGGGGGCTGCCATCACGCCTACATCGTGCTCTGCATTGAGAACAGCTTCATGACT 540
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Prats, Herve
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                                                                      AAAGCTGCGAGCTGCCACAACAGCTACATCGTCCTGTGCATTGAGAATAGCTTCATGACC 543
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Pred. No. 3.1e-76;
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RESULT 9
US-08-985-526-37
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US-08-985-526-37
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Best Local S
Matches 443
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ZIP: 19899
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS_DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/985,52
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TITLE OF INVENTION:
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TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE: 16-JUL-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CLASSIFICATION: PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
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STRANDEDNESS: single
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                         CATGGCTCGGACCCCAACGGCGCAGGCTGACCGAGAGCTACTGTGAGACGTGGCGGACG 420
 CACGGCTCGGACCCCAGTGGGCGGAGGCTGATGGAGAGTTACTGTGAGACATGGCGAACT 430
                                                                    TTTTCTTTTGACGGCAGAGATGTCCTGAGACACCCAGCCTGGCCGCAGAAGAGCCGTATGG
                                                                                                      TTCTCCTTTGACGGCAAGGACGTCCTGAGGCACCCCACCTGGCCCCAGAAGAGCGTGTGG
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Pred. No. 3.1e-76;
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, P.O. Box 2207
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US-09-561-500-12
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SEQ ID NO 12
LENGTH: 573
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Best Local Similarity
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APPLICANT: Philip E. Thorpe
APPLICANT: Rolf A. Brekken
TITLE OF INVENTION: ANTIBODY COMPOSITIONS FOR SELECTIVELY INHIBITING VEGF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches 443;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILE REFERENCE: 4001.002500
CURRENT APPLICATION NUMBER: US/09/561,500
CURRENT FILING DATE: 2000-04-28
PRIOR APPLICATION NUMBER: 60/131,432
PRIOR FILING DATE: 1999-04-28
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OTHER INFORMATION: Description of Artificial Sequence:
OTHER INFORMATION: OLIGONUCLEOTIDE
NAME/KEY: CDS
LOCATION: (1)..(573)
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421 GAGGCTCCCTCGGCCACGGGCCAGGCCTCCTCGCTGCTGGGGGGCAGGCTCCTGGGGCAG
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                                                           CATGGCTCGGACCCCAACGGGCGCAGGCTGACCGAGAGGCTACTGTGAGACGTGGCGGACG
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                                                                                                                                                                           CCCAGCTGGGACTCCCTGTTTTCTGGCTCCAGGGTCAACTGCAACCCGGGGCCCGCATC
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Pred. No. 3.2e-76;
0; Mismatches 103;
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US-09-561-108-12
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TENGTH: 573
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APPLICANT: Rolf A. Brekken

TITLE OF INVENTIONS: ANTHEODY CONJUGATE COMPOSITIONS FOR SELECTIVELY INHIBITING VEGF
FILE REFERENCE: 4001.002584

CUMRENT APPLICATION NUMBER: US/09/561,108

CUMRENT FILING DATE: 1000-04-28

PRIOR APPLICATION NUMBER: 60/131,432

PRIOR APPLICATION NUMBER: 60/131,432

PRIOR PILING DATE: 1999-04-28

NUMBER OF SEQ ID NOS: 44

NUMBER OF SEQ ID NOS: 44

NUMBER OF SEQ ID NOS: 44

SOFTWARE: Patentin Ver: 2.0
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TYPE: DNA
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                         CATGGCTCGGACCCCAACGGGCGAGGCTGACCGAGAGCTACTGTGAGACGTGGCGGACG
                                                                                                                                     TTTTCTTTTGACGGCAGAGATGTCCTGAGACACCCAGCCTGGCCGCAGAAGAGCGTATGG
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Pred. No. 3.2e-76;
0; Mismatches 103;
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RESULT 12
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LENGTH: 573
TYPE: DNA
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Best Local
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CURRENT APPLICATION NUMBER: US/09/561,526

CURRENT FILING DATE: 2000-04-28

PRIOR APPLICATION NUMBER: 60/131,432

PRIOR FILING DATE: 1999-04-28

NUMBER OF SEQ ID NOS: 44

SOFTWARE: Patentin Ver. 2.0
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APPLICANT: Rolf A. Brekken
TITLE OF INVENTION: ANTIBODY CONJUGATE KITS FOR SELECTIVELY INHIBITING VEGF
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LOCATION: (1)..(573)
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AGTGCCGCGAGCTGCCATCACGCCTACATCGTGCTCTGCATTGAGAACAGCTTCATGACT
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Pred. No. 3.2e-76;
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US-09-561-499-12
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SEQ ID NO 12
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APPLICANT: Rolf A. Brekken
TITLE OF INVENTION: ANTIBODY METHODS FOR SELECTIVELY INHIBITING VEGF
FILE REFERENCE: 4001.002582
CURRENT APPLICATION NUMBER: US/09/561,499
CURRENT FILING DATE: 2000-04-28
PRIOR APPLICATION NUMBER: 60/131,432
PRIOR FILING DATE: 1999-04-28
NUMBER: OF SEQ ID NOS: 44
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Pred. No. 3.2e-76;
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US-09-998-831-12
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No. 667694
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APPLICANT: Philip B. Thorpe
APPLICANT: ROLf A. Brekken
IITLE OF INVENTION: ANTIBODY CONJUGATE COMPOSITIONS
TITLE OF INVENTION: INHIBITING VEGF
FILE REFERENCE: 4001.002584
CURRENT APPLICATION NUMBER: US/09/998,831
CURRENT FILING DATE: 2001-11-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT FILING DATE: 2001-11-30
PRIOR APPLICATION NUMBER: 09/561,108
PRIOR FILING DATE: 2000-04-28
NUMBER OF SEQ ID NOS: 44
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FEATURE:
OTHER INFORMATION: Description of Artificial Sequence:
OTHER INFORMATION: OLIGONUCLEOTIDE
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TYPE: DNA
ORGANISM: Artificial Sequence
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   AAAGCTGCGAGCTGCCACAACAGCTACATCGTCCTGTGCATTGAGAATAGCTTCATGACC
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Pred. No. 3.2e-76;
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US-09-231-077D-5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: 60/075,587
PRIOR FILING DATE: 1998-02-23
NUMBER OF SEQ ID NOS: 13
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ORGANISM: human
FEATURE:
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## ALIGNMENTS

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	COMMENT	TITLE JOURNAL	AUTHORS	REFERENCE			ORGANISM	SOURCE	KEYWORDS	VERSION	ACCESSION	DEFINITION	POCUS	BE908201	RESULT 1
Tissue Procurement: ATCC CDNA Library Preparation: Life Technologies, Inc. cDNA Library Preparation: Life Technologies, Inc. cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov plate: LLAM9704 row: d column: 16 High quality sequence stop: 688.	Contact: Robert Strausberg, Ph.D. Email: cqapbs-r@mail.nih.gov	Unpublished (1999)	NIH-MGC http://mgc.nci.nih.gov/.	1 (bases 1 to 707)	Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.	Eukarvota: Metazoa: Chordata: Craniata: Vertebrata: Euteleostomi;	Homo sapiens	Homo sapiens (human)	EST.	BE908201.1 GI:10402537	BE908201	mRNA sequence.	iens cDNA clone IMAGI	mrna linear EST 20-OCT-2000	

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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 757)
                                                     Homo sapiens
                                                                                               mRNA sequence.
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BE906253.1 GI:10399595
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/note="Organ: pancreas; Vector: pCMV-SPORT6; Site_1: Not1; Site_2: Sall; Cloned unidirectionally. Primer: Oligo dT. Average insert size 1.1 kb. Library constructed by Life Technologies."
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/db_xref="taxon:9606"
/clone="IMAGE:3902175"
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/lab_host="DH10B (phage-resistant)"
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Mismatches 0;
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National Institutes of Health, Mammalian
Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
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Tissue Procurement: ATCC
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                                                                     AGTGCCGCGAGCTGCCATCACGCCTACATCGTGCTCTGCATTGAGAACAGCTTCATGACT
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/tissue_type="epithelioid carcinoma"
/lab_host=="DH10B (phage-resistant)"
/clone_lib="NIH_MGC_70"
/note="Organ: pancreas; Vector: pCMV-SPORT6; Site_1: Not1; Site_2: Sall; Cloned unidirectionally. Primer: Oligo dT. Average insert size 1.1 kb. Library constructed by Life Technologies."
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/db_xref="taxon:9606"
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Pred. No. 8.9e-85;
Mismatches 0;
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01 TTCTCCTTTGACGGCAAGGACGTCCTGAGGCACCCGACCTGGCCCCAGAAGAGCGTGTGG 360	41 CCCAGCTGGGAGGCTCTGTTCTCAGGCTCTGAGGGTCCGCTGAAGCCCGGGGCACGCATC 300	181 GTGGGCGTGCCGACCGGCGAGCCGTGCCCATCGTCAACCTCAAGGACGACGAGCTGCTGTTT 240	121 GGGCTGGCGGCACCTTCCGCGCCTTCCTGTCCTCGCGCCTGCAGGACCTGTACAGCAIC 180	61 GGCGGCATGCGGGGGCATCCGCGGGGCCGACTTCCAGTGCTTCCAGCAGGCGCGGGCCGTG 120	1 CACAGCCACCGCGACTTCCAGCCGGTGCTCCACCTGGTTGCGCTCAACAGCCCCCTGTCA 60	y Match 100.0%; Score 546; DB 14; Length 881; Local Similarity 100.0%; Pred. No. 9.3e-85; hes 546; Conservative 0; Mismatches 0; Indels 0; Gaps 0;	National Institutes of Health, Mammalian Gene Collection (MGC) Unpublished (1999) Contact: Robert Strausberg, Ph.D. Email: cgapbs-r@mail.nih.gov Tissue Procurement: Dr. Michael Brownstein cDNA Library Preparation: Invitrogen Corp cDNA Library Preparation: MGC clone distribution: MGC clone distribution: MGC clone distribution: MGC clone distribution information can be found through the I M.A.G.E. Consortium/LLNL at: http://image.llnl.gov Plate: NDAM424 row: g column: 24 High quality sequence stop: 689. Location/Qualifiers 1881 /organism="Homo sapiens" /db xref="taxon:9606" /clome="MMAGE:30356881" /tissue_type="mitutary" /lab host="DHIOB-Ton A ( T1 and T5 phage resistances)" /clome impact	CD105862 CD105862 AGENCOURT 14021788 NIH MGC 179 Homo sapiens cDNA clone IMAGE:30365831 5', mRNA sequence. CD105862 CD105862.1 GI:30759036 EST. Homo sapiens (human) Homo sapiens Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eutleleostomi; Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 881)

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Klausner, R.D., Collins, F.S., Wagner, L., Shenmen, C.M., Schuler, G.D.,

Altschul, S.F., Zeeberg, B., Buetow, K.H., Schaefer, C.F., Bhat, N.K.,

Hopkins, R.F., Jordan, H., Moore, T., Max, S.I., Wang, J., Hsieh, F.,

Diatchenko, L., Marusina, K., Farmer, A.A., Rubin, G.M., Hong, L.,

Stapleton, M., Soares, M.B., Bonaldo, M.F., Casavant, T.L.,

Scheetz, T.E., Brownstein, M.J., Usdin, T.B., Toshiyuki, S.,

Carninci, P., Prange, C., Raha, S.S., Loquellano, N.A., Peters, G.J.,

Abramson, R.D., Mullahy, S.J., Bosak, S.A., McEwan, P.J.,

McKernan, K.J., Malek, J.A., Gunaratne, P.H., Richards, S.,

Worley, K.C., Hale, S., Garcia, A.M., Gay, L.J., Hulyk, S.W.,

Villalon, D.K., Muzny, D.M., Sodergren, E.J., Lu, X., Gibbs, R.A.,

Fahey, J., Helton, E., Ketteman, M., Madan, A., Rodrigues, S.,

Sanchez, A., Whiting, M., Madan, A., Young, A.C., Shevchenko, Y.,

Boutfard, G.G., Blakesley, R.W., Touchman, J.W., Green, E.D.,

Dickson, M.C., Rodriguez, A.C., Schmutz, J., Myers, R.M.,

Schnerch, A., Schein, J.E., Jones, S.J. and Marra, M.A.

Generation and initial analysis of more than 15,000 full-length
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info@bcgsc.bc.ca

Steven Jones, Jennifer Asano, Tan Bosdet, Yaron Butterfield,

Susanna Chan, Readman Chiu, Chris Fjell, Brin Garland, Ran Guin,

Letticia Hsiao, Marrin Krzywinski, Reta Kutsche, Oliver Lee, Soo

Sen Lee, Victor Ling, Carrie Mathewson, Candice McLeavy, Steven

Ness, Pawan Pandoh, Anna-Liisa Prabhu, Parvaneh Saeedi, Jacqueline
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Direct Submission
Submitted (08-DEC-2003) National Institutes of Health, Mammalian Gene Collection (MCC), Cancer Genomics Office, National Cancer Institute, 31 Center Drive, Room 11A03, Bethesda, MD 20892-2590,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BC063833 4230 bp mRNA linear HTC 09-DEC Homo sapiens cDNA clone IMAGE:6181818, containing frame-shift
                                                                                                                                                                                                      cDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Genome Sequence Centre,
BC Cancer Agency, Vancouver, BC, Canada
                                                                                                                                                                                                                                                                                                                                                                                                     NIH-MGC Project URL: http://mgc.nci.nih.gov
Contact: MGC help desk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   human and mouse cDNA sequences
Proc. Natl. Acad. Sci. U.S.A. 99 (26), 16899-16903 (2002)
12477932
                                                                                                                                                                                                                                                                                                                                        Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Dr. James R. Lupski
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Strausberg, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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AGENCOURT 10434549 NIH MGC 109 Homo sapiens cDNA clone IMAGE: 6650260 5', mRNA sequence.
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                                                   BU859398
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This clone was selected for full length sequencing because it passed the following selection criteria: matched mRNA g1: 18765747
This clone has the following problem: frame shifted.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Clone distribution:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
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                                                                                                                                                                                                                            546
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /tissue_type="Peripheral Nervous System, dorsal root
ganglion"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            note="Vector: pCMV-SPORT6"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       clone="IMAGE:6181818"
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                                                944 bp
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
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Contact: Robert Strausberg, Ph.D.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 944)
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National Institutes of Health, Mammalian Gene Collection (MGC)
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                              GCATGGCTCGGACCCCAACGGGCGCAGGCTGACCGAGAGCTACTGTGAGACGTGGCGGAC 293
                                                             GCATGGCTCGGACCCCAACGGGCGCGAGGCTGACCGAGAGCTTGTTGAGACGTTGGCGGAC 419
                                                                                                                           CTTCTCCTTTGACGGCAAGGACGTCCTGAGGCACCCCACCTGGCCCCAGAAGAGCGTGTG
                                                                                                                                                       CTTCTCCTTTGACGGCAAGGACGTCCTGAGGCACCCCACCTGGCCCCAGAAGAGCGTGTG
                                                                                                                                                                                                                            TCCCAGCTGGGAGGCTCTGTTCTCAGGCTCTGAGGGTCCGCTGAAGCCCCGGGGCACGCAT
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                                                                                                                                                                                                                                                                                                                              CGTGCGCCGTGCCGACCGCGCAGCCGTGCCCATCGTCAACCTCAAGGACGAGCTGCTGTT
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                                                                                                                                                                                                                                                                                                                                                                                                                             GGGGCTGGCGGCCTTCCGCGCCTTCCTGTCCTCGCGCCTGCAGGACCTGTACAGCAT
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/lab_host="PHIOB (phage-resistant)"
/clone_lib="NIH_MGC_109"
/note="Organ: ovary; Vector: pOTB7; Site_
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
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Pred. No. 4.5e-80;
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NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Contact: Robert Strausberg, Ph.D. Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   High quality sequence stop: 588.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Unpublished (1999)
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BQ672290
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /tissue_type="epidermoid carcinoma, cell line"
/lab_bost="DH10B (phage-resistant)"
/clone_lib="NHH_MGC_102"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
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97.8%;
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                                                                                                                                                                                                                                                                                                                                                   Email: cgapbs-r@mail.nih.gov
Tissue Procurement: James Martin
CDNA Library preparation: Dr. M. Bento Soares, University of Iowa
cDNA Library Arrayed by: Dr. M. Bento Soares, University of Iowa
cDNA Sequencing by: Dr. M. Bento Soares, University of Iowa
Clone Distribution: Clone distribution information can be obtained
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     181 GTGCGCCGTGCCGACCGCGCAGCCGTGCCCATCGTCAACCTCAAGGACGAGCTGCTGTTT 240
                                                                                                                                                                                                                                                                                         from Dr. M. Bento Soares, bento-soares@wiowa.edu
The following repetitive elements were found in this cDNa
The following repetitive elements were found in this cDNa
Sequence: 1-44, >POLY_A#Simple_repeat (matched compliment)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          241
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            347 GGGCTGGCGGGCATCCTTCCGCGCCTTCCTGTCCTCGCGCCTGCAGGACCTGTACAGCATC 406
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                         POLYA=Yes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Tumor Gene Index
Unpublished (1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BU615520 703 bp mRNA
UI-H-FG0-bcp-p-10-0-UI.s1 NCI CGAP EN1 2
UI-H-FG0-bcp-p-10-0-UI 3', mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens (human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BU615520.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AGAGTG-CCGCGAGCTGCCATCACGCCTACATCGTGCTCTGCATTGAGAACAGCTTCATG 537
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GAGGCTCCCTCGGCCÁCGGCCAGGCCTCCTTCCCTGGGGGGGGGAGGCTCCCTGGGGC 706
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GAGGCTCCCTCGGCCACGGGCCAGGCCTCCTCGCTGCTGGGGGG--CAGGCTCCTGGGGGC 478
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CATGGCTCGGACCCCAACGGGCGCAGGCTGACCGAGAGCTACTGTGAGACGTGCCGGACG 646
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CATGGCTCGGACCCCAACGGCCCAGGCTGACCGAGAGCTACTGTGAGACGTGGCGGACG
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                                                                                                                                                                                                                                                                                       primer: M13 FORWARD
/clone lib="NCI_CGAP_ENI_2"
/note="Organ: Bone; Vector: pT7T3-Pac (Pharmacia) with a modified polylinker; Site 1: EcoR I; Site_2: Not I; NCI_CGAP_ENI_2 is a cDNA Tibrary containing the following
                                                                                                                        /organism="Homo sapiens"
/mol_type="mkNA"
/db_xref="kaxon:9606"
/clone="UI-H-FGO-bcp-p-10-0-UI"
/tissue_type="Enchondroma cell line"
                                                                                 /dev_stage="Adult"
/lab_host="DH10B (Life Technologies)"
                                                                                                                                                                                                                                                 Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GI:23281735
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   _2 Homo sapiens cDNA clone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Anatomy Project (CGAP),
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Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     702 GGTGCGCTCAACAGCCCCCTGTCAGGCGGCATGCGGGGCATCCGCGGGGCCGACTTCCAG
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                                                                                                                                                                                                                                                                                                                    682 bp mRNA linear EST 15-OCT-19: xd20f03.xl NCI CGAP Ov23 Homo sapiens cDNA clone IMAGE:2594333 3' similar to SW:CA1H HUMAN P39060 COLLAGEN ALPHA 1(XVIII) CHAIN [CONTAINS: ENDOSTATIN] ;, mRNA sequence.
NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap
                                    Mammalia; Eutheria;
1 (bases 1 to 682)
                                                                                                          Eukaryota; Metazoa;
                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                      AW089583.1 GI:6046927
                                                                                                                                                                                                                                                                                                       AW089583
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ACCTGGCCCCAGAAGACGTGTGGCATGGCTCGGACCCCAACGGGCGCAGGCTGACCGAG 396
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AACCTCAAGGACGAGCTGCTGTTTCCCAGCTGGGAGGCTCTGTTCTCAGGCTCTGAGGGT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        library was constructed according to Bonaldo, Lennon and Soares, Genome Research, 6:791-806, 1996. First strand containing a Not I site. Double stranded cDNA was ligated to an EccR I adaptor, digested with Not I, and cloned directionally into pT/T3-Pac vector. The oligonucleotide used to prime the synthesis of first-strand cDNA contains a library tag sequence that is located between the Not I site and the (GT)18 tail. The sequence tag for this site and the (GT)18 tail. The sequence tag for this lames Martin from University of lowa.

TAG_TISSUE-Enchondroma cell line (Mix of EN1 and EN2)
                                                                                                                                                                                        sapiens (human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        tissue(s): Enchondroma cell line (2 cell lines). The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TAG_LIB=UI-H-FG0
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99.2%;
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                                                                              Chordata; Craniata; Vertebrata; Euteleostomi; primates; Catarrhini; Hominidae; Homo.
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Pred. No. 1.5e-77;
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cDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Center
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consorting/LLNL at:
                                                                                                                                                                                                                                                                      460
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                                    202
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Emmert-Buck, M.D., Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Contact: Robert Strausberg,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Unpublished (1997)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         www-bio.llnl.gov/bbrp/image/image.html
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                                ATTGAGAACAGCTTCATGACTGCCTCC 176
                                                                                              ATTGAGAACAGCTTCATGACTGCCTCC 546
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/note="Ovgan: ovary; Vector: pCMV-SPORT6; Site 1: Sall;
Site_2: NotI; Cloned unidirectionally. Primer: Oligo dT.
Average insert size 1.35 kb. Tumor types include: mixed
Mullerian tumor, papillary serous, clear cell, spindle
cell. All are primary tumors, metastasis positive. Life
Technologies catalog #: 11534-013"
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(lab_host="DH10B"
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/db_xref="taxon:9606"
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Pred. No. 5.8e-76;
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                        Query Match 89.7%;
Best Local Similarity 99.8%;
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Genetics (www.resgen.com).

The following repetitive elements were found in this cDNA seguence: 1-44, >POLY_A#Simple_repeat (matched compliment)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Tissue Procurement: Dr. Gregg Hageman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Coordinated Laboratory for Computational Genomics
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CDNA Library preparation: Dr. M. Bento Soares, Univeristy of Iowa CDNA Library Arrayed by: Dr. M. Bento Soares, Univeristy of Iowa DNA Sequencing by: Dr. M. Bento Soares, Univeristy of Iowa Clone Distribution: Researchers may obtain clones from Research
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            University of Iowa
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97044477
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Mammalla; Butheria; Primates; Catarrhini; Hominidae; Homo.
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          Conservative
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: 319 335 9565
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            bento-soares@uiowa.edu
                                                                                                                                                                       the synthesis of first-strand CDNA contains a library tag sequence that is located between the Not I site and the (dT).18 tail. The sequence tag for this library is CGCGTNATACC. This library was created for the program, Gene Discovery in the Visual System, supported by National Eye
                                                                                                                                                                                                                                                                                                                                                  /lab_host="DH10B (Life Technologies) (T1 phage resistant)" (clone_lib="UI-E-E01"  
/note="Organ: eye; Vector: pT7T3-Pac (Pharmacia) with a modified polylinker; Site 1: EcoR I; Site 2: Not I; UI-E-E01 is a normalized cDNA library containing the following tissue(s): fetal eye. The library was constructed according to Bonaldo, Lennon and Soares, Genome Research, 6:791-806, 1996. First strand cDNA  
Genome Research, 6:791-806, 1996. First strand cDNA
                                                                                                                                                                                                                                                                   synthesis was primed with an oligo-dT primer containing a Not I site. Double stranded cDNA was ligated to an EcoR I adaptor, digested with Not I, and cloned directionally into pT7T3-Pac vector. The oligonucleotide used to prime
                                                                                                                        TAG_TISSUE=human fetal eye
TAG_LIB=UI-E-EO1
                                                                                                        TAG_SEQ=CGCGTATACC"
                                                                                                                                                                 Institute (NEI)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        tissue_type="fetal eye"
|dev_stage="fetal"
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    Score 490; DB 12;
Pred. No. 4.3e-75;
0; Mismatches 1;
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296 GCATCTTCTCCTTTGACGGCAAGGACGTCCTGAGGCACCCCACCTGGCCCAGAAGAGGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     563 GCATCGTGCGCCGTGCCGACCGCGCAGCCGTGCCCATCGTCAACCTCAAGGACGAGCTGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      176 GCATCGTGCGCCGACCGCCGCAGCCGTGCCCATCGTCAACCTCAAGGACGAGCTGC
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                                                                                                                                                                                                                  CDNA Library Preparation: Ling Hong/Rubin Laboratory CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: NIH Intramural Sequencing Center Clone distribution: MGC clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov plate: LLCM1306 row: a column: 14 High quality sequence stop: 679.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            56 TGTCAGGCGGCATGCGGGGCATCCGCGGGGCCGACTTCCAGTGCTTCCAGCAGGCGCGGG 115
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; I
Mammalia; Butheria; Primates; Catarrhini; Hominidae;
1 (bases 1 to 832)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         Email:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NIH-MGC http://mgc.nci.nih.gov/
National Institutes of Health,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 cgapbs-r@mail.nih.gov
/tissue_type="adenocarcinoma cell line"
/lab_host="DH10B (phage-resistant)"
/clone_lib="NIH_MGC_15"
                                                                      /mol_type="mRNA"
/db_xref="taxon:9606"
/clone="IMAGE:4582933"
                                                                                                                                                 organism="Homo sapiens"
                                                                                                                                                                                                  location/Qualifiers
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274 GGCGGCATGCGGGGCATCCGCGGGGCCGACTTCCAGTGCTTCCAGCAGGAGGCGGGCCGTG
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                                                                                                                                                                                                                                                                                        979 bp mRNA
AGENCOURT 8414390 NIH_MGC_102 Homo sapiens
5', mRNA sequence.
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                                                                    Contact: Robert Strausberg, Ph.
Email: cgapbs-r@mail.nih.gov
                                                                                                                         NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene
                                                                                                                                                           Mammalia; Eutheria; Primates; 1 (bases 1 to 979)
                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                Homo sapiens
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BQ673186.1 GI:21784020
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                                                    Tissue Procurement: ATCC
                                                                                                        Unpublished (1999)
                                                                                                                                                                                                                            Homo sapiens (human)
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cDNA Library Preparation: Rubin Laboratory cDNA Library Arrayed by: The I.M.A.G.E. Consortium DNA Sequencing by: Agencourt Bioscience Corporation
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97.6%;
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574 361 514 301 454 241

VERSION KEYWORDS

SOURCE

ORGANISM

COMMENT

sapiens cDNA clone IMAGE:5019339

linear

EST 05-JUL-2001

REFERENCE

AUTHORS TITLE JOURNAL

LOCUS BQ673186 RESULT 11

DEFINITION ACCESSION

В Ş B S 밁

693 479 634 419 밁 8 밁

61

ORIGIN

Matches

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Local

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334

121

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Clone distribution: MGC clone distribution information found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Plate: LLCM2450 row: g column:
High quality sequence stop: 638.
                                                                  /note="Organ: salivary gland; Vector: pOTB7; Site_1: XhoI; Site_2: EcoRI; cDNA made by oligo-dT priming. Birectionally cloned into EcoRI/XhoI sites using the following 5' adaptor: GGCACGAG(G). Library constructed by Ling Hong in the laboratory of Gerald M. Rubin (University of California, Berkeley) using ZAP-cDNA synthesis kit (Stratagene) and Superscript II RT (Life Technologies). Note: this is a NIH_MGC Library."
                                                                                                                                                                                                                                                                                                       /tissue_type="epidermoid carcinoma, cell line"
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/clone_lib="NHH_MGC_102"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                  clone="IMAGE:6272287"
                                                                                                                                                                                                                                                                                                                                                                                                                          /mol_type="mRNA"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       organism="Homo sapiens"
DΒ
13;
Length
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CACAGCCACCGCGACTTCCAGCCGGTGCTCCACCTGGTTGCGCTCAACAGCCCCCCTGTCA
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                                                                                                                                                                                                                                                                                                                                                                                                                                    GTGCGCCGTGCCGACCGCGCAGCCGTGCCCATCGTCAACCTCAAGGACGAGCTGCTGTTT
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                                                                                                                 GAGGCTCCCTCGGCCACGGCCACGGCCTCCTCGCTG--CTGGGGGGCAGGCTCCTGGGGG
                                                                                                                                                                       CATGGCTCGGACCCCAACGGCGCAGGCTGACCGAGAGCTACTGTGAGACGTGGCCGACG
                                                                                                                                                                                                                CATGGCTCGGACCCCAACGGGCGCAGGCTGACCGAGAGCTACTGTGAGACGTGGCGGACG
                                                                                                                                                                                                                                                          TTCTCCTTTAACGGCAAGGACGTCCTGAGGCACCCCACCTGGCCCCAGAAGAGCGTGTGG
                                                                                                                                                                                                                                                                                          TTCTCCTTTGACGGCAAGGACGTCCTGAGGCACCCCACCTGGCCCCAGAAGAGCGTGTGG
                                                                                                                                                                                                                                                                                                                                            CCCAGCTGGGAGGCTCTGTTCTCAGGCTCTGAGGGTCCGCTGAAGCCCCGGGGCACGCATC
<u>AGAGTGCCCGCGAGCTGCCATCACGCCCTACATCGTGGCTCTGCATTGAGAACAGCTTCA</u>
                             AGAGTG-CCGCGAGCTGCCATCACG-CCTACATCGT-GCTCTGCATTGAGAACAGCTTCA 535
                                                                                          ĊĠĠĠĊĊĀĠĠĊĊŤĊĊŤĊĠĊŤĠĠĊŢĠĠĠĠĠĠĊĀĀĠĊŤĊĊĊŢĠĠĠĠĊ
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Pred. No. 3.2e-74;
0; Mismatches 8
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mRNA sequence

EST 16-JAN-2002

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191 CCGACCGCGCAGCCGTGCCCATCGTCAACCTCAAGGACGAGCTGCTGTTTCCCAAGCTGGG 250
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  131 GCACCTTCCGCGCCTTCCTGTCCTCGCGCCTGCAGGACCTGTACAGCATCGTGCGCCGTG 190
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                High quality sequence stop: 769.
                ACGGCAAGGACGTCCTGAGGCACCCCACCTGGCCCCAGAAGAGCGTGTGGCATGGCTCGG 370
                                                                                                                                                                                                                                                                                                         AGGCTCTGTTCTCAGGCTCTGAGGGTCCGCTGAAGGCCCGGGGCACGCATCTTCTCCTTTG 310
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 (bases 1 to 884)
NHH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           EST
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BI161007.1 GI:14621008
                                                                                         ACGGCAAGGACGTCCTGAGGCACCCACCTGGCCCCAGAAGAGGCGTGTGGCATGGCTCGG
                                                                                                                                                                                                                                                                      AGGCTCTGTTCTCAGGCTCTGAGGGTCCGCTGAAGCCCGGGGCACGCATCTTCTCCTTTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                cDNA Library Preparation: Ling Hong/Rubin Laboratory cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Incyte Genomics, Inc. Clone distribution information can be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens (human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GGGGCATCCGCGGGCCGACTTCCAGTGCTTCCAGCAGGCGCGGGCCGTGGGGGCTGGCGG 61
                                                                                                                                                                                                                                                                                                                                                             CCGACCGCGCAGCCGTGCCCATCGTCAACCTCAAGGACGAGCTGCTGTTTCCCCAGCTGGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                    GCACCTTCCGCGCCTTCCTCTCGCGCGCCTGCAGGACCTGTACAGCATCGTGCGCCGTG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /note=Torgan: pancreas; Vector: pOTB7; Site 1: XhoI; Site 2: EGORI; cDNA made by oligo-dT priming.
Directionally cloned into EccRI/XhoI sites using the following 5' adaptor: GGCAGCAGG(0). Size-selected >500bp for average insert size 1.8kb. Library constructed by Ling Hong in the laboratory of Gerald M. Rubin (University of California, Berkeley) using ZAP-cDNA synthesis kit (Stratagene) and Superscript II RT (Life Technologies).
Note: this is a NIH_MGC Library. | "
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /tissue_type="epithelioid carcinoma cell line"
/lab_host="DH10B (phage-resistant)"
/clone_lib="NIH_MGC_42"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /mol_type="mRNA"
/db_xref="taxon:9606"
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100.0%; Prr
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Pred. No. 1.2e-72;
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AV696242
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Best Local Similarity
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MEDLINE
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240 TCCCAGCTGGGAGGCTCTGTTCTCAGGCTCTGAGGGTCCGCTGAAGCCCGGGGCACGCAT
                                                                                                                                                             186 GGGCTGGCGGCACCTTACGCGCCCTTTCTGTCCTCGCGCCCTGCAGGACACTGTACAGCAT
                                                                                                    180 CGTGCGCCGTGCCGACCGCGCAGCCGTGCCCATCGTCAACCTCAAGGACGAGCTGCTGTT 239
                                                                                                                                                                                                121 GGGCTGGCGGGCACCTTCCGCGCCCTTCCTCGCGCCCTGCAGGAC-CTGTACAGCAT 179
                                                                                                                                                                                                                                                      126 GGCGGCATGCGGGGCATTCGCGGGGCCGACTTCCAGTGCTTTCAGCAGGCGCGGGCCGTG
                                                                                                                                                                                                                                                                                                                                                                                                                                               508;
                                                                                                                                                                                                                                                                                         61 GGCGGCAIGCGGGGGCAICCGCGGGGCCCGACTICCAGCTGCTTCCAGCAGGCGCGGGCCGTG
                                                                                                                                                                                                                                                                                                                                            66 CACAGTCACCGCGACTTCCAGCCGGTGCTCCACCTGTTTGCGCTCAACAGACCCCCTGTCA 125
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   422 GCTGCCATCACGCCTACATCGTGCTCTGCATTGAGAACAGCTTCATGACTGCCTCC 477
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               491 GCTGCCATCACGCCTACATCGTGCTCTGCATTGAGAACAGCTTCATGACTGCCTCC 546
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                                                                                                                                                                                                                                                                                                                                                                                1 CACAGCCACCGCGACTTCCAGCCGGTGCTCCACCTGGTTGCGCTCAACAGCCCCCTGTCA 60
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Insight into hepatocellular carcinogenesis at transcriptome level by comparing gene expression profiles of hepatocellular carcinoma with those of corresponding noncancerous liver Proc. Natl. Acad. Sci. U.S.A. 98 (26), 15089-15094 (2001)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        בוואבייג Contact: Zeguang Han
Contact: Zeguang Han
Chinese National Human Genome Center at Shanghai
ייי Shouiing Road, Zhangjiang Hi-Tech Park, Pudong,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Email: hanzg@chgc.sh.cn
This clone is available at CHGC in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AV696242
AV696242 GKC Homo sapiens cDNA clone GKCEAG02 5', AV696242
AV696242.1 GI:10298105
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                                                                      CGTGCGCGTGCCGACCGCGAACCGTGCCCATCGTCAACCTCAAGGACCAGCTGGTGTT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /tissue_type="hepatocellular carcinoma"
/dev_stage="Adult"
/lab_host="SOLR"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /mol_type="mRNA"
/db_xref="taxon:9606"
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92.9%;
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Pred. No. 2.3e-72;
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Shanghai

FEATURES

source

COMMENT

REFERENCE

AUTHORS TITLE

SOURCE KEYWORDS VERSION ACCESSION

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Matches 476; Query Match

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AUTHORS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BQ723254 1093 bp mRNA linear BST 16-JUL-200: AGENCOURT 8486770 Lupski_sympathetic_trunk Homo sapiens cDNA clone IMAGE:6196366 5', mRNA sequence.
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BQ723254.1 GI:21862151
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Dr. James R. Lupski
CDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
CDNA Sequencing by: Agencourt Bioscience Corporation
DNA Sequencing by: Agencourt Bioscience Corporation
Clone distribution: MGC clone distribution information can
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
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Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                http://image.lln1.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Contact: Robert Strausberg, Ph.D
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Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      through the I.M.A.G.E. Consortium/LLNL at:
                                            5'-GACTACTTCTAGATCGCGAGCGGCCCT(15)-3'. Size selected > 1 kb for average insert length 1.9 kb. This is a primary library, non-amplified. Library constructed by Life Technologies and donated by J. Lupski, M.D./Ph.D. (Baylor college of Medicine); available through Life
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      612
                                                                                                                                                              /clone_lib="Lupski_sympathetic_trunk"
/clone_lib="Lupski_sympathetic_trunk"
/note="Wector: pCMV-SPORT6 (Life Technologies); Site_1:
NotI; Site_2: SalI; cDNA made bilgo-dT priming.
Directionally cloned using the following adaptors:
5'-TCGACCCCACGCGTCCG-3' and
                                                                                                                                                                                                                                                                          /tissue_type="sympathetic trunk"
/dev_stage="adult, 16 yr"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                         /organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                      /clone="IMAGE:6196366"
                                                                                                                                                                                                                                                                                                                                               'sex="male"
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BU632049.1
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UI-H-FEI-bea-m-09-0-UI.sl NCI GAP FEI HC
UI-H-FEI-bea-m-09-0-UI 3', mRNA sequence.
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Unpublished (1997)
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1 (bases 1 to 664)
NCI CGAP http://www.ncbi.nlm.nih.gov/ncicgap
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
                                                                                                   from Dr. M. Bento Soares, bento-soares@uiowa.edu
The following repetitive elements were found in
sequence: 1-35, ANT rich#Low_complexity (matched
                                                                                                                                                                                                                                                                      Email: cgapbs-r@mail.nih.gov
Tissue Procurement: James Martin
                                                                                                                                                                                                                                                                                                                Contact: Robert Strausberg, Ph.D
                                                                                                                                                                                                                                                                                                                                                                                National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens (human)
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                                                                                                                                                                       cDNA Library preparation: Dr. M. Bento Soares, University of Iowa cDNA Library Arrayed by: Dr. M. Bento Soares, University of Iowa DNA Sequencing by: Dr. M. Bento Soares, University of Iowa Clone Distribution: Clone distribution information can be obtained Clone Distribution:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CCCAGAAGAGCGTGTGGCATGGCTCGGACCCCAACGGGCGCAGGCTGACCGAGAGCTACT
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/organism="Homo sapiens"
                                        Location/Qualifiers
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Pred. No. 1.8
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NCI CGAP FE1 Homo sapiens cDNA clone
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                                                                                                                   (matched compliment)
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480

420

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Query Match 83.0%;
Best Local Similarity 99.8%;
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244 GCTCTGCATTGAGAACAGCTTCATGACTGCCTCC
                                                             513 GCTCTGCATTGAGAACAGCTTCATGACTGCCTCC 546
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                                                                                                                                                                                             GCTGCTGGGGGCAGGCTCCTGGGGCAGAGTGCCGCGAGCTGCCATCACGCCTACATCGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GGGTCCGCTGAAGCCCCGGGGCACGCATCTTCTCCTTTGACGGCAAGGACGTCCTGAGGCA 332
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TAG_TISSUB-Human grade 2 chondrosarcoma cell line pool TAG_LIB-UI-H-FB1
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/note="Organ: Chondrosarcoma; Vector: pT773-Pac
(Pharmacia) with a modified polylinker; Site 1: Ecor I;
Site 2: Not I; NCI_CGAP_FEI is a normalized cDNA library
derived from a pool of mRNA obtained from 3 cell lines
from grade II chondrosarcoma tissues. The library was
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /clone="UI-H-FE1-bea-m-09-0-UI"
/tissue_type="Cell lines"
/dev_stage="Adult"
/lab_host="DH10B (Life_Technologies)"
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/db_xref="taxon:9606"
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Pred. No. 1.1e-68;
0; Mismatches 1;
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Search completed: September 15, 2004, 20:10:47 Job time : 2457  ${\tt secs}$ 

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Gapop 10.0 , Gapext 1.0
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546
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Copyright (c) 1993 - 2004 Compugen Ltd.
                                                                    CGM2 6/ptodata/2/pubpna/US09E PUBCOMB.seq:*
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//CGM2 6/ptodata/2/pubpna/US09C PUBCOMB.seq:*
//CGM2 6/ptodata/2/pubpna/US09 NEW PUB.seq:*
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//CGM2 6/ptodata/2/pubpna/US10B PUBCOMB.seq:*
//CGM2 6/ptodata/2/pubpna/US10C PUBCOMB.seq:*
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/cgn2_6/ptodata/2/pubpna/US60_PUBCOMB.seq:*
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                      6/ptodata/2/pubpna/US10 NEW PUB.seq:*
6/ptodata/2/pubpna/US60 NEW PUB.seq:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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Result No.	Score	Query Match	Query Match Length DB	; ₩	ID	Description
1	546	100.0	546	15	US-10-042-347-4	Sequence 4, Appli
23	546	100.0	549	12	US-10-135-872B-10	
ω	546	100.0	549	15	US-10-131-241-53	
4	546	100.0	549	15	US-10-292-418-3	
ហ	546	100.0	552	9	US-09-873-676-30	
a	546	100.0	552	13	US-09-978-531-2	
7	546	100.0	552	17	US-10-607-501-1	
89	546	100.0	555	17	US-10-449-609-3	
9	546	100.0	627	13	US-09-373-938-4	
10	546	100.0	1158	17	US-10-449-609-5	
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12	546	100.0	4551	15	US-10-060-036-144	
13	546	100.0	4875	16	US-10-264-049-835	
14	545	99.8	551	14	US-10-080-797-2	

Best Local Similarity
Matches 546; Conserval

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17	17	15	17	16	16	10	13	15	16	15	15	14	13	15						13	13	10	10	15	15	15	15	15	13	17
-10-469-285-4	-10-433-793-3	L	10-433-793-3	US-10-159-563-67	US-10-133-937-67	US-09-971-392-159	占	US-10-060-036-64	10-430-503-3	US-10-131-241-51	1	US-10-080-797-4	US-09-373-938-1	US-10-292-418-17	US-10-373-561-12	US-09-998-831-12	US-10-036-869-37	-60	US-09-775-174-3	US-10-210-172-165	US-10-210-172-163	-09-938-391-	-09-938-391-3	-10-292-418-3	-10-131-241-5		US-10-131-241-59	US-10-042-347-6	US-10-210-172-161	US-10-432-364-4
423,	31	Sequence 45.	Sequence 32,	Sequence 67,	Sequence 67, 1	Sequence	Sequence 145	Sequence 64,	37	5	70	4.	Sequence 1, Appli	17	12	Sequence 12, Appl	O			Sequence 165, App		Ь.		е 34.	50.	48,	OD.	Sequence 6, Appli	Sequence 161, App	Sequence 4, Appli

## ALIGNMENTS

RESULT 1 US-10-042-347-4

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US-10-042-347-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: US 09/315,689
PRIOR FILING DATE: 1999-05-20
PRIOR APPLICATION NUMBER: US 60/106,343
PRIOR FILING DATE: 1998-10-30
PRIOR APPLICATION NUMBER: US 09/154,302
PRIOR APPLICATION NUMBER: US 09/740,168
PRIOR APPLICATION NUMBER: US 08/740,168
PRIOR APPLICATION NUMBER: US 08/740,168
PRIOR APPLICATION NUMBER: US 60/005,835
PRIOR APPLICATION NUMBER: US 60/005,835
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 4, Application US/10042347 Publication No. US20030114370A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                               SEQ ID NO 4
                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: O'Reilly, Michael S.
APPLICANT: Folkman, M. Judah
TITLE OF INVENTION: Nucleic Acid Molecules Encoding Endostatin Protein and Peptide
TITLE OF INVENTION: Thereof
                                                                                                                                                                                                                                                                        NUMBER OF SEQ ID NOS: 6
SOFTWARE: PatentIn version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILE REFERENCE: 05213-0880 (43170-249874)
CURRENT APPLICATION NUMBER: US/10/042,347
CURRENT FILING DATE: 2002-01-11
                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION NUMBER: US 60/026,263 PRIOR FILING DATE: 1996-09-17
                                                                                                                                               TYPE: DNA ORGANISM: Homo sapiens
                                                                                                                                                                                                             LENGTH: 546
                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US 60/023,070 FILING DATE: 1996-08-02
                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: 1995-10-23
100.0%; Score 546; DB 15; ilarity 100.0%; Pred. No. 1.9e-132; Conservative 0; Mismatches 0;
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US-10-135-872B-10
                                                       US-10-135-872B-10
                                                                                                                      PRIOR APPLICATION NUMBER: US 60/287,673
PRIOR FILING DATE: 2001-04-30
PRIOR APPLICATION NUMBER: US 60/370,634
PRIOR FILING DATE: 2002-04-05
NUMBER OF SEQ ID NOS: 13
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 10, Application US/10135872B Publication No. US20040071659A1
Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                 APPLICANT: McArthur, James
APPLICANT: Patel, Salil
APPLICANT: Patel, Salil
APPLICANT: Jooss Karin
APPLICANT: Mendez, Michael
APPLICANT: Donahue, Brian
ITILE OF INVENTION: Viral-Mediated Delivery and In Vivo Expression of
ITILE OF INVENTION: Polymicleotides Encoding Anti-Angiogenic Proteins
FILE REFERENCE: 3802-709-27
                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Chang, APPLICANT: Wu, We
                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/10/135,872B CURRENT FILING DATE: 2002-04-29
                                                                       TYPE: DNA
ORGANISM: Homo
                                                                                                           LENGTH: 549
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APPLICANT: Holaday, John W.
APPLICANT: Fortier, Anne H.
TITLE OF INVENTION: Compositions and Methods for Inhibiting Endothelial Cell Prolifera
TITLE OF INVENTION: And Regulating Angiogenesis Using Cancer Markers
FILE REFERENCE: 05213-0344 43170-271565
FILE REFERENCE: 05213-0344 43170-271565
FILE REFERENCE: 05213-0344 43170-271565
FILE REFERENCE: 05213-0344 43170-271565
FILE REFERENCE: 05213-0344 43170-27156
FILING DATE: 1902-07-22
PRIOR APPLICATION NUMBER: US 09/413,049
PRIOR APPLICATION NUMBER: US 09/413,049
PRIOR APPLICATION NUMBER: US 09/316,802
PRIOR FILING DATE: 1999-05-21
PRIOR FILING DATE: 1999-05-22
                                                                                                                  SEQ ID NO 53
                                                                                                                                                                               NUMBER OF SEQ ID NOS: 65
                                                                                                                                                    SOFTWARE: PatentIn
TYPE: DNA
ORGANISM: Homo
                                                                            LENGTH:
                                                                        549
       sapiens
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Query Match Best Local Similarity Matches 546; Conserv

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Score 546; DB 15; Pred. No. 1.9e-132;

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Sequence 53, Application US/10131241 Publication No. US20030012792A1

GENERAL INFORMATION:

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; FEATURE:
NAME/KEY: CDS
LOCATION: (1)..(549)
COTHER INFORMATION: endostatin
US-10-292-418-3
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US-10-292-418-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; Sequence 3, Application US/10292418 ; Publication No. US20030139365A1 ; GENERAL INFORMATION:
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     Query Match
Best Local Similarity
Matches 546; Conserv
                                                                                                                                                                                                            SOFTWARE: PatentIn Ver. 2.0 SEQ ID NO 3
                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Lo, Kin-Ming
APPLICANT: Li, Yue
APPLICANT: Gillies, Stephen D
TITLE OF INVENTION: Expression and Export of Angiogenesis Inhibitors
TITLE OF INVENTION: Immunofusins
                                                                                                                                                                                                                                            CURRENT APPLICATION NUMBER: US/10/292,418
CURRENT FILING DATE: 2002-11-12
PRIOR APPLICATION UNMBER: 09/383,315
PRIOR FILING DATE: 1999-08-25
PRIOR PELICATION NUMBER: US 60/097,883
PRIOR FILING DATE: 1998-08-25
NUMBER OF SEQ ID NOS: 54
                                                                                                                                                                                                                                                                                                                                                                     FILE REFERENCE: LEX-006C1
                                                                                                                                                      LENGTH: 549
TYPE: DNA
ORGANISM: Homo sapiens
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 100.0%; Score 546; DB 15; ilarity 100.0%; Pred. No. 1.9e-132; Conservative 0; Mismatches 0;
                                  Length 549;
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SEQ ID NO 30
LENGTH: 552
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                                                                                                                 Query Match
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APPLICANT: Sim, Kim L.
TITLE OF SIM, Kim L.
TITLE REFERENCE: 05213-0378 (43170-25933)
CURRENT APPLICATION NUMBER: US/09/873,676
CURRENT FILING DATE: 2001-06-04
PRIOR FILING DATE: 2000-06-02
PRIOR FILING DATE: 2000-06-02
                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: US 60/289,387 PRIOR FILING DATE: 2001-05-08
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                                                                              100.0%; Score 546; DB 9; 1 ilarity 100.0%; Pred. No. 1.9e-132; Conservative 0; Mismatches 0;
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; LENGTH: 552
; TYPE: DNA
; ORGANISM: Homo :
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                                                                                                                                                                                                                     Matches
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Best Local S
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APPLICANT: Liang, Hong
TITLE OF INVENTION: Angiogenesis-Inhibiting Peptides and Proteins and Methods of Use
FILE REFERENCE: 05213-0811 43170-264992
CURRENT APPLICATION NUMBER: US/09/978,531
CURRENT FILING DATE: 2001-10-15
                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: US 60/240,127
PRIOR FILING DATE: 2000-10-13
NUMBER OF ESE ID NOS: 13
SOFTWARE: PatentIn version 3.1
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US-10-607-501-1

Sequence 1, Application US/10607501

Publication No. US20040091465A1

PUBLICATION:
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                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Yim, Zachary
ITIE OF INVENTION: Therapeutic Antiangiogenic Compositions and Methods
FILE REFERENCE: 05213-0791 (43170-286879)
CURRENT APPLICATION NUMBER: US/10/607,501
CURRENT ELING DATE: 2003-06-26
CURRENT FILING DATE: 2003-06-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: DNA
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                                                    CCCAGCTGGGAGGCTCTGTTCTCAGGCTCTGAGGGTCCGCTGAAGCCCCGGGGCACGCATC
                                                                                             GTGCGCCGTGCCGACCGCGCAGCCGTGCCCATCGTCAACCTCAAGGACGAGCTGCTGTTT
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Pred. No. 1.9e-132;
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publication No. US20040127407A1
; GENERAL INFORMATION:
APPLICANT: CHEN, WEN Y
TITLE OF INVENTION: HUMAN PROLACTIN ANTAGONIST-ANGIOGENESIS INHIBITOR
TITLE OF INVENTION: FUSION PROTEINS
FILE REFERENCE: 035879-0163
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-10-449-609-3
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Best Local Similarity
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CURRENT FILING DATE: 2003-09-26
PRIOR APPLICATION UNMBER: 60/384,121
PRIOR FILING DATE: 2002-05-31
NUMBER OF SEQ ID NOS: 45
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ORGANISM: Homo sapiens
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100.0%; Score 546; DB 13;
Best Local Similarity 100.0%; Pred. No. 1.8e-132;
Matches 546; Conservative 0; Mismatches 0;
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/09/373,938
CURRENT FILING DATE: 1999-08-13
NUMBER OF SEQ ID NOS: 17
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APPLICANT: Chen, Cheauyun Theresa
TITLE OF INVENTION: ADENOVIRAL VECTORS INCLUDING DNA SEQUENCES ENCODING ANGIOGENIC
FILE REFERENCE: 4-3089991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME/KEY: CDS
LOCATION: (1)..(627)
OTHER INFORMATION:
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ORGANISM: Homo sapiens
FEATURE:
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                                CATGGCTCGGACCCCAACGGCGCAGGCTGACCGAGAGCTACTGTGAGACGTGGCGGACG
                                                                                                                                                     CATGGCTCGGACCCCAACGGGCGCAGGCTGACCGAGAGCTACTGTGAGACGTGGCGGACG
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GAGGCTCCCTCGGCCACGGCCAGGCCTCCTCGCTGCTGGGGGGGCAGGCTCCTGGGGCAG
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RESULT 10 US-10-449-609-5

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Sequence 5, Application US/10449609
Publication No. US20040127407A1
GENERAL INFORMATION:
APPLICANT: CHEN, WEN Y
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OTHER INFORMATION: Description of Artificial Sequence: OTHER INFORMATION: hPRL_G129R-Endostatin Fusion nucleotide sequence
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AGTGCCGCGAGCTGCCATCACGCCTACATCGTGCTCTGCATTGAGAACAGCTTCATGACT 1146
                         AGTGCCGCGAGCTGCCATCACGCCTACATCGTGCTCTGCATTGAGAACAGCTTCATGACT 540
                                                                              GAGGCTCCCTCGGCCACGGCCAGGCCTCCTCGCTGCTGGGGGGCAGGCTCCTGGGGGCAG
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NUMBER OF SEQ ID NOS: 3950
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 2178
LENGTH: 3394
TYPE: DNA
ORGANISM: Homo sapiens
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US-09-880-107-2178
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Patent No. US20020142981A1
GENERAL INFORMATION:
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CURRENT APPLICATION NUMBER: US/09/880,107
CURRENT FILING DATE: 2001-06-14
PRIOR APPLICATION NUMBER: US 60/211,379
PRIOR FILING DATE: 2000-06-14
PRIOR APPLICATION NUMBER: US 60/237,054
PRIOR APPLICATION NUMBER: US 60/237,054
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TITLE OF INVENTION: Gene Expression Profiles in Liver Cancer
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Scherf, Uwe
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                                                             AGTGCCGCGAGCTGCCATCACGCCTACATCGTGCTCTGCATTGAGAACAGCTTCATGACT
                                                                                                                  GAGGETECETEGGECACGGECEAGGECTECTEGETGGTGGGGGGCAGGETECETGGGGCAG
                                    AGTGCCGCGAGCTGCCATCACGCCTACATCGTGCTCTGCATTGAGAACAGCTTCATGACT
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US-10-449-609-5

Matches 546; Query Match Best Local

h 100.0%; Similarity 100.0%;

Conservative

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SOFTWARE: PatentIn Ver. 2.1 SEQ ID NO 5

LENGTH: 1158 TYPE: DNA

ORGANISM: Artificial Sequence

FEATURE:

PRIOR FILING DATE: 2002-05-31 NUMBER OF SEQ ID NOS: 45

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APPLICANT: Benson, Darin R.

APPLICANT: Kalos, Michael D.

APPLICANT: Lodes, Michael D.

APPLICANT: Persing, David H.

APPLICANT: Persing, David H.

APPLICANT: Hepler, William T.

APPLICANT: Jiang, Yugiu

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY

TITLE OF INVENTION: AND DIAGNOSIS OF PANCREATIC CANCER

FILE REFERENCE: 210121.566

CURRENT APPLICATION NUMBER: US/10/060,036

CURRENT FILING DATE: 2002-01-30

NUMBER OF SEQ ID NOS: 4560

SOPTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 144

LENGTH: 4551
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US-10-060-036-144
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                                                         AGTGCCGCGAGCTGCCATCACGCCTACATCGTGCTCTGCATTGAGAACAGCTTCATGACT 4539
                                                                                                                                               GAGGCTCCCTCGGCCACGGCCAGGCCTCCTCGCTGCTGGGGGGCAGGCTCCTGGGGGAAG 480
                                 GCCTCC 546
                                                                              AGTGCCGCGAGCTGCCATCACGCCTACATCGTGCTTCTGCATTGAGAACAGCTTTCATGACT 540
                                                                                                                               GAGGCTCCCTCGGCCA
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Pred. No. 1.6e-132;
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US-10-264-049-835
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Best Local Similarity
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PRIOR FILING DATE: 2000-06-07
NUMBER OF SEQ ID NOS: 4360
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CURRENT FILING DATE: 2002-10-04
PRIOR APPLICATION NUMBER: PCT/US01/18569
PRIOR FILING DATE: 2001-06-07
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TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
FILE REFERENCE: PA133P1
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ORGANISM: Homo
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                                  541 GCCTCC 546
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RESULT 14
US-10-080-797-2
is Sequence 2, Application US/10080797
publication No. US20020183253A1
GENERAL INFORMATION:

OCULAR

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Sequence 4, Application US/10432364

publication No. US20040091996A1

GENERAL INFORMATION:
APPLICANT: VIRGENS HOTECHNOLOGY LIMITED
APPLICANT: VIRGENS HOTECHNOLOGY LIMITED
TITLE OF INVENTION: A VIRUS WHICH CAN EXPRESS TUP
TITLE OF INVENTION: SPECIFIC TUMOR CELLS AND THE
FILE REFERENCE: iecolog2pct
CURRENT APPLICATION NUMBER: US/10/432,364
CURRENT FILING DATE: 2003-11-06
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; ORGANISM: Human
US-10-080-797-2
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APPLICANT: Dixon, Katharine H.
APPLICANT: Brazzell, Romulus K.
TITLE OF INVENTION: METHOD FOR TREATING OCUL
TITLE OF INVENTION: METHOD FOR TREATING OCUL
FILE REFERENCE: 4-31891A
FULE REFERENCE: 4-31891A
CURRENT APPLICATION NUMBER: US/10/080,797
CURRENT FILING DATE: 2002-02-21
NUMBER OF SEQ ID NOS: 21
NUMBER OF SEQ ID NOS: 21
SOFTWARE: FastSEQ for Windows Version 4.0
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SEQ ID NOS: 36
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Search completed: September 15, 2004, 21:20:46 Job time : 357 secs

OM nucleic Run on:

nucleic search, using sw model

GenCore version 5.1.6 Copyright (c) 1993 - 2004 Compugen Ltd.

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Minimum
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Perfect score:
Sequence:
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Pred. No. is the number of results predicted by chance to have a

score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

## Result No. 418 381.2 381.2 381.2 381.2 381.2 381.2 381.2 381.2 381.2 381.2 381.2 381.2 381.2 520 426.2 419.6 418 418 418 544.4 544.4 538 534 534 Score 1100.0 100.0 10 100.0 Query Match 100.0 100.0 100.0 100.0 100.0 100.0 Length DB AF257775 BD102800 BD128313 AF018081 AF416592 BC037060 AR194060 AR193166 AR193166 AX100092 AX370823 BD272245 BD272245 BD240900 AX399631 BD188110 E34073 BD271283 AR183471 AR183520 AR217279 AX816108 I51045 AX409531 HUMCOL18AX AF018082 AR268609 AR100642 AR282751 AX042272 AK130835 AK098216 AX473835 BD081407 AR431862 AF282883 AX370851 IJ AX816106 AK098216 Homo sapi AK816106 Sequence AK816108 Sequence AK816108 Sequence 151045 Sequence AK409531 Sequence 122548 Human colla AF018082 Homo sapi AF018081 Homo sapi AF018081 Homo sapi AF184060 Homo sapi AF184060 Homo sapi AF184060 Homo sapi AF189166 Sequence AX100092 Sequence ED272245 Anti-angi BD272245 Anti-angi BD272246 Anti-a AR217279 Sequence AR282751 Sequence AX042272 Sequence AX490666 Sequence AX473835 Sequence BD081407 Fused pro AR431862 Sequence AF282883 Homo sapi AX370851 Sequence AK130835 Homo sapi BD266741 Proteins AX100086 Sequence Description

## ALIGNMENTS

SOURCE SOURCE ORGANISM REFERENCE AUTHORS TITLE JOURNAL FEATURES	RESULT 1 AR193165 LOCUS DEFINITION ACCESSION VERSION VERSION
Unknown. Unknown. Unknown. Unclassified. 1 (bases 1 to 546) 0'Reilly,M.S. and Folkman,M.Judah. Therapeutic antiangiogenic endostatin compositions Patent: US 6346510-A 4 12-FEB-2002; Location/Qualifiers	AR193165 546 bp DNA linear PAT 20-APR-2002 Sequence 4 from patent US 6346510. AR193165 AR193165.1 GI:20239130

source

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BD240882.1 GI:33050652
JP 2002523036-A/2.
                                                                                                                                                                    Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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      Homo sapiens (human)

JP 2002523036-A/2
30-UUL-2002
35-AUG-1999 JP 2000566305
25-AUG-1998 US 60/097883
KIN MING LO, YUE LI, STEPHEN D GILLIES
C12N15/09, A61K31/711, A61K38/00, A61K39/395, A61K48/00, A61P7/00,
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/mol_type="unassigned
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AX100086
AX100086.1 GI:13539061
Liang, H., Sim, K.L., Chang-Murad, A., Zhou, X., Madsen, J. Boerner, R.J., Bermejo, L.L., Mistry, F.R., Shepard, S.R.
                                            Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                      Homo sapiens
                                                                                    Homo sapiens
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A61P43/00,C07K14/745,C07K14/78,C07K19/00,C12N5/10,C12P21/02,
C12N15/00,
C12N5/00,A61K37/02
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Location/Qualifiers
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
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                                                                                    (human)
                                                                                                                                          549 bp
Patent W00119989
                                            Chordata; Craniata; Vertebrata; Euteleostomi; Primates; Catarrhini; Hominidae; Homo.
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Pred. No. 1.2e-77;
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RESULT 4
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KEYWORDS
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                                                                                                                  SOURCE
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 Eukaryota; Metazoa; Chordata; Craniata; Ver
Mammalla; Butheria; Primates; Catarrhini; H
1 (bases 1 to 552)
Macdonald,N.J. and Sim,K.L.
Proteins that bind anglogenesis-inhibiting
and methods of use thereof
Patent: JP 2002532068-A 2 02-OCT-2002;
                                                                                                                           BD266741
BD266741.1 GI:33076509
JP 2002532068-A/2.
                                                                                                                                                                  Proteins that bind angiogenesis-inhibiting proteins, compositions and methods of use thereof.
                                                                                                  Homo sapiens
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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Pred. No. 1.2e-77;
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                                                                     Catarrhini; Hominidae; Homo.
                                                                                  Craniata; Vertebrata; Euteleostomi;
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Query Match
Best Local Similarity
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OS Homo sap
PN JP 20025
PD 02-OCT-2
PF 06-DEC-1
PR 04-DEC-1
PI NICHOLAS
PC C12N15/0
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C07K14/705,G01N33/15,G01N33/50,G01N33/53,G01N33/566//A61K35/76,
GCCTCC 546
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                                                                         AGTGCCGCGAGCTGCCATCACGCCTACATCGTGCTCTGCATTGAGAAACAGCTTCATGACT
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JP 2002532068-A/2
02-OCT-2002
06-DEC-1999 JP 2000585271
04-DEC-1998 US 09/206059
NICHOLAS J MACDONALD, KIM LEE SIM
C12N15/09, A61K38/00, A61P9/00, A61P27/02, A61P29/00, A61P35/00,
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
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Sequence 2 from Patent WO0230982.
AX473835
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/mol_type="unassigned DNi
/db_xref="taxon:9606"
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Eukaryota; Metazoa;
Mammalia; Eutheria;
                                     BD081407.1 GI:22627010
JP 2001518304-A/50.
Homo sapiens (human)
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Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Chordata; Primates;

Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.

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30-SEP-1998 JP 2000513958
01-OCT-1997 US 60/060609
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BARBARA K KLEIN, JOHN P MCKEARN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MARK A BOLANOWSKI, MAIRE H CAPARON, GERALD F CASPERSON, SUSAN
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JP 2001518304-A/50
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
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Location/Qualifiers
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Pred. No. 1.2e-77;
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Homo sapiens multi-functional protein MFP mRNA, complete cds
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Violand, B.N. and Harding, E.I.
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Direct Submission
Submitted (27-JUN-2000) R & D, Rc
Submitted (27-JUN-2000) R & D, Rc
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Mammalia; Eutheria; Primates;
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                                                                                      AGTGCCGCGAGCTGCCATCACGCCTACATCGTGCTCTGCATTGAGAACAGCTTCATGACT
                                                                                                            AGTGCCGCGAGCTGCCATCACGCCTACATCGTGCTCTGCATTGAGAACAGCTTCATGACT
                                                                                                                                                                      GAGGCTCCCTCGGCCACGGCCAGGCCTCCTCGCTGCTGGGGGGCAGGCTCCTGGGGCAG
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/protein id="AAM52249.1"
/protein id="AAM52249.1"
/db_xref="gi:121425751"
/db_xref="gi
    780
                                           546
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/mol_type="mRNA"
/db_xref="taxon:9606"
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Pred. No. 1.1e-77;
Nismatches 0;
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1, MI 48239, USA
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/mol_type="unassigned NNA"
/db_xref="taxon:32630"
/note="mdNA for mIgSP fused to (pEnd-HR#1"
                                                                                                 882
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576 240 516 180 456 120 396 60

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REFERENCE AUTHORS

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Submitted (31-JUL-2003) Sumio Sugano, Institute of Medical Science, University of Tokyo, Laboratory of Genome Structure, Human Genome Center; Shirokane-dai, 4-6-1, Minato-ku, Tokyo 108-8639, Japan (E-mail:flcdna@ims.u-tokyo.ac.jp, Tel:81-3-5449-5286,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           full insert sequencing: RAB and Helix Research Institute.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Fax:81-3-5449-5416)
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Homo sapiens cDNA FLJ27325 fis, clone TWS08226, highly similar
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/clone_lib="TMS"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /mol_type="mRNA"
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/clone="TMS08226"
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                                                                              1644 CACAGCCACCGCGACTTCCAGCGGTGCTCCACCTGGTTGCGCTCAACAGCCCCCTGTCA 1703
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61 GGCGGCATGCGGGGCATCCGCGGGGCCGACTTCCAGTGCTTCCAGCAGCAGGCGCGGGCCCGTG
                                                                                                                    1 CACAGCCACCGCCGACTTCCAGCCGGTGCTCCACCTGGTTGCGCTCAACAGCCCCCCTGTCA 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Submitted (04-JUL-2002) Takao Isogai, FLJ Project(HRI Team); 2-6-7 Kazusa-Kamatari, Kisarazu, Chiba 292-0812, Japan (E-mail-genomicsehri.co.jp, Tel:81-438-52-3975, Fax:81-438-52-3986) MEDO human cDNA sequencing project supported by Ministry of Economy, Trade and Industry of Japan; cDNA full insert sequencing: Research Association for Biotechnology (Rab); cDNA library construction: Helix Research Institute (HRI) (supported by Japan Key Technology Center etc.); 5'- & 3'-end one pass sequencing: RAB, HRI, and Biotechnology Center, National Institute of Technology and Evaluation; clone selection for full insert sequencing: HRI and RAB; annotation: HRI and RAB.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Kawai-Hio,Y., Saito,K., Nishikawa,T., Kimura,K., Yamashita,H., Matsuo,K., Nakamura,Y., Sekine,M., Kikuchi,H., Kanda,K., Wagatsuma,M., Murakawa,K., Kanehori,K., Sugiyama,A., Rawakami,B., Suzuki,Y., Sugano,S., Nagahari,K., Masuho,Y., Nagai,K. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AK098216 2364 bp mRNA linear Homo sapiens cDNA FLJ40897 fis, clone UTERUZ002964, to Homo sapiens type XVIII collagen (COL18A1) mRNA.
                                                                                                                                                                                                                   Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NEDO human cDNA sequencing project
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Arita, M., Musashino, K., Yuuki, H., Hara, H., Sugiyama, T., Irie, R., Otsuki, T., Sato, H., Wakamatsu, A., Ishii, S., Yamamoto, J., Isono, Y.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Oshima, A., Takahashi-Fujii, A., Tanase, T., Imose, N.,
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                                                                                                                                                                                                                                                                                                                                                /tissue_type="uterus"
/clone_lib="UTERU2"
                                                                                                                                                                                                                                                                                                                    note="cloning"
                                                                                                                                                                                                                                                                                                                                                                                                                           /mol_type="mRNA"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            organism="Homo sapiens"
                                                                                                                                                                                                                                                                                                                                                                                                      clone="UTERU2002964"
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Pred. No. 8.1e-78;
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                                                                                                                                                                                                                                      DB 9; Length 2364;
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Takeuchi, K.,

highly similar PRI 15-JUL-2002

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Qy  1 CACAGCCACCGCGACTTCCGCGGGCCCTTCCTGCCGCGGGGCCGGACCTGTACAGCATC 180  Qy  1 CACAGCCATCGCGGACTTCCAGCCGGGCCTCCACCAGGACCTGTACAGCACCCTGTCA 132  Db  73 CACAGCCACCGCGACTTCCAGCCGGGGCCGACTTCCAGCAGGGCGGGGCCGTGTCA 132  Qy  61 GGCGGCATGCGGGGATTCCGGGGGGCCGACTTCCAGTGTTTCCAGCAGGGGGGGG	synthetic construct  Maynthetic construct  Maynthetic construct  Autificial sequences.  E 1  S Mertins, P., Celik, I., Kisker, O., Sleep, D., Hay, J. and Albumin-fused anti-angiogenesis peptides  Aventis Behring GmbH (DE) , Delta Biotechnology Limi Location/Qualifiers  1. 2376  /organism="synthetic construct" /mol type="unassigned DNA" /db_xref="taxon:32630"  /note="Description of Artificial Sequence: the N-terminal endostatin-albumin fusion op frame"	2 SULT 13 816106 CUS CESSION FRION WORDS	Db 1704 GGCGGCATGCGGGGGCCTTCCGGGGGCCGACTTCCAGTGCTTCCAGGCAGG
Qy  1 CACAGCCACCGCGACTTCCAGCCGGTGCTACCACCTGGTACACCCCCCTGTCA 60	AUTHORS Merrins, P., Celik, I., Kisker, O., Sleep, D., Hay, J. and Hauser, H.P. AUTHORS Merrins, P., Celik, I., Kisker, O., Sleep, D., Hay, J. and Hauser, H.P. Albumin-fused anti-angiogenesis peptides JOURNAL Patent: WO 03066085 A 32 14-NUG-2003; Aventis Behring GmbH (DE) J. APUG-2003; Aventis Behring GmbH (DE) J. Delta Biotechnology Limited (GB) Location/Qualifiers  1. 2376 //onl type="unassigned DNA" //db_xref="taxon:32630" //nol type="unassigned DNA" //db_xref="taxon:32630" //nol type="unassigned DNA" //db_xref="taxon:32630" //nol type="unassigned DNA" //db_xref="taxon:32630" //nol type="unassigned DNA" //onl type="unassigned DNA" //on	Oy 541 GCCTCC 546	193 GAGGCTGGC 181 GTGGGCC 241 CCCAGCT 241 CCCAGCT 301 TTCTCCT 373 TTCTCCT 373 TTCTCCT 421 GAGGCTC 421 GAGGCTC 421 GAGGCTC 421 GAGGCTC 421 GAGGCTC 433 GAGGCTC 433 GAGGCTC 434 AGTGGCTC 435 AGTGGCTC 437 AGTGGCTC 437 AGTGGCTC 438 AGTGGCTC 438 AGTGGCTC 439 GAGGCTC 441 AGTGGCTC 441 A

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2044 GCCTCC 2049	541 GCCTCC 546	AGTGCCGCGAGCTGCCATCACGCCTACATCGTGCTCTGCATTGAGAACAGCTTCATGACT 2043	481 AGTGCCGCGAGCTGCCATCACGCCTACATCGTGCTCTCCATTGAGAACAGCTTCATGACT 540	GAGGCTCCCTCGGCCACGGCCCAGGCCTCCTCGCTGCTGGGGGGCAGGCTCCTGGGGGCAG	GAGGCTCCCTCGGCCACGGGCCAGGCCTCCTCGCT	CATGGCTCGGACCCCAACGGCGCAGGCTGACCGAGAGCTACTGTGAGACGTGGCGGACG 1923	CATGGCTCGGACCCCAACGGCCGCAGGCTGACCGAGAGCTACTGTGAGACGTGGCGGACG 420	TTOTCCTTTGACGCAAGGACGTCCTGAGGCACCCCACCTGGCCCCAGAAGAGCGTGTGG 1863	TICTCTTTGACGGCAAGGACGTCCTGAGGCACCCCACCTGGCCCCCAGAAGAGCGTGTGG 360	

Search completed: September 15, 2004, 19:29:43 Job time : 2359 secs

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Database :
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 200000000
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546
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Copyright (c) 1993 - 2004 Compugen Ltd.
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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Abv94763 Human pan	Himat	_	Aax78379 Human alp	Himar	Ξ.		-	DNA er	Human	Himar		cDNA 6	-	Nuclea	Human e	SEO II				Aaz51291 Human ang	Aax77719 Human end	Additional end		Description		

Claim 10; Page 39; 68pp; English.

angiogenesis, useful birth control agents.

P-PSDB; AAY94323.

Novel endostatin capable of inhibiting endothelial cell proliferation and angiogenesis, useful for treating angiogenesis-dependent cancers and as

<b>4</b> 5	44	4.3	4.2	4 0	40	. L.	000	37	6	i Cr	34	. L	32	<u>د</u> د د	3 0	29	28	2/	0 0	) (	J I	44
381.2	381.2	381.2	381.2	405.6	416.4	418	418	419.6	426.2	520	532.6	534	534	538	542.8	542.8	542.8	542.8	045	ı U	7 (	546
69.8	69.8	69.8	69.8	74.3	76.3	76.6	76.6	76.8	78.1	95.2	97.5	97.8	97.8	98.5	99.4	99.4	99.4	99.4	99.8	· C		100.0
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AAD18701	AAC88290	AAX84635	AAX79949	ADE40259	ADE40257	AAL46062	AAL46063	AAZ51309	AAZ50398	ABK09978	AAA64014	AAS00868	AAA27005	ADE40255	AAL44000	ABQ76079	AAL55454	AAH79104	ABQ81193	AAX77720	CCERCAGA	プロ / O F F
					Human N	Canine	Canine				Aaa64014 Nucleotid		•	•		-	Speci	Human		Aax77720 Human col	Abq54955 Human ova	

## ALIGNMENTS

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RESULT 1
AAA27004
                                                                                  O'reilly MS, Folkman MJ;
                                                                                                                                                                                                                                                                                     Human; endothelial cell proliferation inhibitor; collagen XVIII; angiogenesis inhibitor; anti-tumour; cytostatic; antipsoriatic; vasotropic; determatological; ophthalmological; vulnerary; antiarteriosclerotic; antidiabetic; haemostatic; contraceptive; coular angiogenic disease; atherosclerosis; scleroderma; myocardial angiogenesis; telangiectasia; angiofibroma;
                                                                                                                         30-OCT-1998;
20-MAY-1999;
                                                                                                                                                     01-NOV-1999;
                                                                                                                                                                                           WO200026368-A2.
                                                                                                                                                                                                                                     Key
CDS
                                                                                                     (CHIL-) CHILDRENS MEDICAL CENT
                                                                                                                                                                          11-MAY-2000.
                                                                                                                                                                                                                                                               Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                               Human endostatin gene.
                                                                                                                                                                                                                                                                                                                                                                                                       AAA27004;
                                                                                                                                                                                                                                                                                                                                                                                                                        AAA27004 standard; DNA; 546 BP.
                                                                                                                                                                                                                                                                                  wound granulation,
                                                                                                                                                                                                                                                                                                                                                                                   11-AUG-2000
                                                               2000-365617/31.
                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                        98US-0106343P.
99US-00315689.
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                                                                                                                                                                                                                        /*tag=
                                                                                                                                                                                                                                           Location/Qualifiers
                                                                                                                                                                                                             /product= "Endostatin protein"
                                                                                                                                                                                                                                                                                   ds.
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10-AUG-1999

(first entry)

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cc lung carcinomas. There was tumour mass regression non-detectable levels cafter 12 days of therapy due to the angiogenesis inhibitory activity of ce endostatin. Thus the protein is useful for treatment of angiogenesis. Cc endostatin are useful for treating and diagnosis of tumours, ocular cangiogenic disease, Osler-Webber syndrome, myocardial angiogenesis, cc angiogenic disease, Osler-Webber syndrome, myocardial angiogenesis, cc plaque neovascularisation, telangiectasia, haemophiliac joints, cangiofibroma and wound granulation, for treatment of diseases related to excessive or abnormal stimulation of endothelial cells e.g. intestinal cc adhesions, atherosclerosis, scleroderma. The protein may also be useful cc as a birth control agent by reducing or preventing uterine cc vascularisation. The gene for endostatin may be isolated from cells or tissue that express high levels of endostatin, eg. tumour cells, by chemical country and servence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present sequence encodes an endostatin protein which is the carboxy terminal protein of human collagen XVIII. Recombinant mouse endostatin (20 mg/kg) was administered subcutaneously to mice implanted with Lewis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local
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Pred. No. 5.1e-102;
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anti-angiogenic; attenuation; cytostatic; anti-diabetic; ophthalmology;
tumour growth; solid tumour; diabetic retinopathy; retina; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               inhibiting fusions and fragments, where the varial vector is attenuated for use in human gene therapy. The products of the invention have anti-angiogenic, cytostatic, anti-diabetic and ophthalmological activity. The vector is used in gene therapy for inhibiting tumour growth in humans harbouring a solid tumour. The vector expresses an anti-angiogenic polypeptide. An additional use comprises treatment of diabetic angiogenic polypeptide. An anti-angiogenic polypeptide inhibits angiogenesis
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      retinopathy, where the anti-angiogenic polypeptide inhibits angiogenesis in the vicinity of the retina. The vector is administered to cells ex vivo and then administered to the patient
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         This invention describes a novel viral gene therapy vector comprising a nucleic acid molecule encoding an anti-angiogenic polypeptide chosen from human or murine angiostatin, human or murine endostatin and angiogenesis inhibiting fusions and fragments, where the viral vector is sufficiently
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Pred. No. 5.1e-102,
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                                                                                                                                                                                                                                       Novel fusion protein of angiostatin or endostatin and an immunoglobulin FC region, useful for treating conditions mediated by angiogenesis, sucl as rheumatoid arthritis, tumors and macular degeneration.
                                                                                                                                                                                                                                                                                                                                                                                                                                               Lo K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  metastasis; atherosclerosis; psoriasis; rheumatoid arthritis; ocular angiogenic disease; diabetic retinopathy; macular degeneration; myocardial angiogenesis; plaque neovascularisation; telangiectasia;
                                                                                                                                                                                                                                                                                                                                                                 P-PSDB;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; immunoglobulin gamma Pc fragment; endostatin; immunofusin; angiogenesis; inhibitor; cytostatic; antirheumatoid; antiarthritic; antipsoriatic; antidiabetic; ophthalmological; immunoguppressant;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     vasotropic; vulnerary; treatment; antiarteriosclerosis;
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a signal sequence, an immunoglobulin Fc region, and an angiogenesis inhibitor selected from angiostatin, endostatin, a plasminogen fragment having angiostatin activity, a collagen XVIII fragment having endostatin activity, or combinations of them. The fusion protein (immunofusin) is used to inhibit angiogenesis and to treat diseases or conditions mediated

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AAA29884 standard; cDNA; 549

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22-AUG-2000 AAA29884;

(first entry)

Human angiogenesis inhibiting factor 1 encoding cDNA.

by angiogenesis.

Conditions that may be treated

include

The patent discloses a DNA molecule encoding a fusion protein comprising

Claim 12; Page 40-41; 68pp; English.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              blood born tumours, tumour metastasis, benign tumours including haemangiomas, acoustic neuromas, neurofibromas, trachomas and premulomas, rheumatoid arthritis, psoriasis, ocular angiogenic of granulomas, rheumatoid arthritis, psoriasis, ocular angiogenic ocuparation arthritis, ocular arthritis, ocular
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abnormal vessel disease; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present sequence encodes an angiogenesis inhibiting factor (I), designated IAF-1. The present invention also describes: (1) preparation of (I) and its derivative; (2) an IAF binding acceptor and its preparation; and (3) an IAF antibody. (I) is useful for preparing new biological preparations for effectively treating various tumours and abnormal-vessel diseases. The IAF antibody is preferably a polyclonal antibody, mosaic antibody, single stranded antibody and human originated
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                                                                                                             TTCTCCTTTGACGGCAAGGACGTCCTGAGGCACCCACCTGGCCCCAGAAGAGCGTGTGG
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No. 5.1e-102;
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WO200119989-A2

22-MAR-2001

14-SEP-2000; 2000WO-US025166

99US-0153698P

(ENTR-) ENTREMED INC

Liang H, S: Bermejo LL, Sim KL, Mistry Chang-Murad A, y FR, Shepard SR, Zhou Schrimsher × Madsen en J, JL; Boerner RJ;

2001-244802/25.

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541 GCCTCC 546

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P-PSDB; AAU00896, AAU00897, AAU00898, AAU00899
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Producing Endostatin protein for treating angiogenesis mediated diseases such as solid tumors, comprises recombinantly producing the protein using an expression system, and recovering and purifying the protein.

Claim 6; Page 29; 67pp; English

e.g., diabetic retinopathy, retinopathy of prematurity, macular degeneration, corneal graft rejection, neovascular glaucoma, colon cancer, retrolental fibroplasia, rubeosis, Osler-Webber Syndrome, myocardial angiogenesis, plaque neovascularisation, telangiectasia, haemophiliac joints, angiofibroma, and wound granulation. Endostatin(TM) is also useful for treating disease of excessive or abnormal stimulation of endothelial cells such as intestinal adhesions, atherosclerosis, scleroderma and hypertrophic scars. Higher yields of more purified, and biologically active Endostatin(TM) are obtained by the new method. Endostatin(TM) can be stored in buffers for extended periods of time, and also subjected to lyophilisation, while preserving biological activity. Centrifugation of broth from fermentation steps in production is avoided, preventing unwanted potential cellular lypis and contamination with additional proteins, pigments, enzymes and other cellular chemicals and granulomas, rheumatoid arthritis, psoriagis, ocular angiogenic diseases invention is useful for producing, recovering and purifying Endostatin (TM) from biological sources, such as biological fluids, tissues, cell culture media, and fermentation media. Endostatin (TM) is useful for haemangioma, acoustic neuromas, neurofibromas, trachomas, and pyogenic treating angiogenesis mediated diseases such as solid tumours, blood The sequence encodes Human Endostatin(TM). The new method of the invention is useful for producing, recovering and purifying Endo orne tumours, leukaemias, tumour metastases, benign tumours e.g.

Sequence 549 BP; 82 A; 196 C; 178 G; 93 T; 0 U; 0 Other;

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Best Local Sim
Matches 546;
                                        Query Match
                             Similarity
                      Conservative
                         100.0%;
                    °.
                          Score 546; DB 5; ]
Pred. No. 5.1e-102;
                    Mismatches
                   <u>.</u>
                                   Length 549;
                  Indels
                 0;
                Gaps
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뮹 Š 밁 Ş 밁 121 121 5 61 GGGCTGGCGGGCACCTTCCGCGCCTTCCTGTCCTCGCGCCTGCAGGACCTGTACAGCATC 180 GGGCTGGCGGCACCTTCCGCGCCTTCCTGTCCTCGCGCCTGCAGGACCTGTACAGCATC 180 GGCGGCÄTGCGGGGGCATCCGCGGGGGCCGACTTCCAGTGCTTCCAGCAGGCGCGGGGCCGTG CACAGCCACCGCGACTTCCAGCCGGTGCTCCACCTGGTTGCGCTCAACAGCCCCCTGTCA CACAGCCACCGCGACTTCCAGCCGGTGCTCCACCTGGTTGCGCTCAACAGCCCCCTGTCA 120 60

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RESULT 6
ABA00774
ID ABAC 밁 07-NOV-2002. WO200288173-A2 Gene; human; plasminogen; angiostatin; neovascularisation; kringle domain; cell proliferation; viral vector; replication-defective; cancer; tumour; ss. Human endostatin coding sequence Homo sapiens. 18-MAR-2003 ABA00774; ABA00774 standard; cDNA; 541 GCCTCC (first entry 546 /product= "Endostatin" Location/Qualifiers /\*tag= L. .549 מ 549

29-APR-2002; 2002WO-US013461

30-APR-2001; 2001US-0287673P. 05-APR-2002; 2002US-0370634P.

(CELL-) CELL GENESYS INC

Wu WW, Macarthur J, Patel S, Jooss K, Mendez

P-PSDB; WPI; 2003-129131/12. AAG79753

New recombinant viral vector expressing human angiostatin useful for inhibiting anglogenesis in a mammalian subject with cancer or tumor.

Example 4; Page 80-82; 83pp; English.

inhibiting angiogenesis in a mammal, especially with cancer or a Note: This sequence is given incorrectly in the sequence listing specification as an amino acid sequence This sequence encodes endostatin. Endostatin is a 20 kD C-terminal fragment of collagen XVIII that inhibits angiogenesis. The endostatin coding sequence may be used in the recombinant viral vector of the invention for obtaining angiostatin activity. The vector comprises a promoter capable of expressing human angiostatin operably linked to a structural gene encoding one or more domains of human angiostatin. The vector, which may be a replication-defective viral vector, is useful for inhibition angiorance; in a mammal consociation with

Sequence 549 BP; 82 A; 196 C; 178 G; 93 T; 0 U; 0 Other:

S Best Loca Matches Query Match Local Similarity Conservative 100.0%; 0; Score 546; DB 7; I Pred. No. 5.1e-102; Mismatches 0 Length 549; Indels 0;

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Mckearn JP;
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  The specification describes multifunctional proteins which comprise combinations of angiostatin, endostatin, interferon, thrombospondin,
                                                                                                                                                   New multifunctional proteins useful
                                                                                 Disclosure;
                                                                                                                                                                                                        WPI; 1999-255098/21
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AAA68203; 27-OCT-2000

(first entry)

AAA68203 standard;

cDNA; 552

Human endostatin cDNA sequence.

Angiogenesis-inhibiting protein receptor; angiogenesis; angiostatin; endostatin; plasminogen; laminin; treatment; wound healing; solid tu psoriasis; scleroderma; myocardial angiogenesis; Crohn's disease;

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Pred. No. 5.1e-102;
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cc similar to that of a plasminogen fragment (see murine plasminogen ABB16490). Angiostatin has the ability to inhibit angiogenesis. Endostatin is also an angiogenesis inhibiting protein (see ABB16451 and CC AAA68203). Sequences AAA68242 and AAB16522 represent coding and protein (see equences of human laminin. Laminin is an angiostatin binding protein, CC and some of the peptides of the invention share homology with regions of CL laminin. Peptides AAB16452-B16521 (excluding AAB16490) are the angiogenesis-inhibiting protein receptor fragments of the invention. The CC peptides bind either angiostatin or endostatin and can be used in methods for treating diseases and processes that are mediated by angiogenesis, crohn's disease, cerebral collaterals, arteriovenous malformations, collicious, diabetic retinopathy, arthritis, wound healing, peptic ulcers,
                                                                                                                                                                                                                                                                                                                                                                                                                      Matches 546;
                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  rubeosis, diabetic retinopathy, arthritis, wound healing, peptic ulcers, Helicobacter related diseases, fractures, placentation and cat scratch fever. They are useful for the detection and prognosis of cancer. DNA sequences A628204-A628241 encode the peptides of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       the DNA sequences encoding them. Angiogenesis is the generation of new blood vessels into a tissue, and normally occurs in wound healing, foetal and embryonal development and the formation of the corpus luteum, endometrium and placenta. Angiostatin is a protein (see AAB16450 and AAA68202) involved in angiogenesis, and has an amino acid sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 552 BP; 83 A; 196 C; 179 G; 94 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New angiogenesis-inhibiting protein receptors, useful in methods for treating diseases and processes that are mediated by angiogenesis, so solid tumors, psoriasis, scleroderma and myocardial angiogenesis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This invention relates to angiogenesis-inhibiting protein receptors,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; Page 86; 100pp; English.
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                                                                  GTGCGCCGTGCCGACCGCAGCCGTGCCCATCGTCAACCTCAAGGACGAGCTGCTGTTT
                                                                                                                                                                                                                                                                      GGCGGCATGCGGGGCATCCGCGGGGGCCGACTTCCAGTGCTTCCAGCAGCAGCGCGGGCCGGGCCGTG 120
                                                                                                                                                                                         GGGCTGGCGGCACCTTCCGCGCCCTTCCTGTCCTCGCGCCTGCAGGACCTGTACAGCATC 180
                                                                                                                                                                                                                                   GGCGGCATGCGGGGCATCCGGGGGGCCGACTTCCAGTGCTTCCAGCAGGCGCGGGGCCGTG 120
                                                                                            GTGCGCCGTGCCGACCGCGCAGCCGTGCCCATCGTCAACCTCAAGGACGAGCTGCTGTTTT
                                                                                                                                                    GGGCTGGCGGGCACCTTCCGGCGCCTTCCTCGCGCGCCTGCAGGACCTGTACAGCATC
                                                                                                                                                                                                                                                                                                                       CACAGCCÁCCGCGÁCTTCCÁGCCGGTGCTCCÁCCTGGTTGCGCTCÁÁCAGCCCCCTGTCÁ 60
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Pred. No. 5.1e-102;
                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
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밁 S 밁 Ś 맑 Ş 뮍 Ş

in Streptomyces. Leader sequences of Streptomyces sp. strain C5 SnpA S. venezuelae alpha-amylase proteins are linked to the N-terminal of endostatin. This ensures that endostatin protein is produced as a secreted, soluble protein which needs no refolding, is stable in the fermentation broth and is produced in large quantities. The method is

The present sequence encodes human endostatin. The protein is expressed in Streptomyces. Leader sequences of Streptomyces sp. strain C5 SnpA and S. venezuelae alpha-amylase proteins are linked to the N-terminal of

used for preparing soluble recombinant human, murine or primate endostatin, which is useful in the treatment of cancer, inhibition of tumour growth, inhibition of angiogenesis, isolation of receptors for

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                                                                                                                        Preparation of soluble recombinant endostatin involves transforming Streptomycete host with expression vector comprising nucleotide sequence encoding endostatin operably linked to linker and leader peptide.
                                                                                                        Example 1; Fig 6; 57pp; English.
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                                                                                                                                                                                                                                                        13-APR-1999;
                                                                                                                                                                                                                                                                           12-APR-2000; 2000WO-US009747
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                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                    Streptomyces sp. strain C5; SnpA; S. venezuelae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Nucleotide sequence of human endostatin encoded by plasmid pMALcH#15.
                                                                                                                                                                                                                                 (MERI ) MERCK & CO INC.
                                                                                                                                                                                                                                                                                                                                                                                                                         endostatin; cancer; tumour growth; angiogenesis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     06-MAR-2001
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                                                                                                                                                                                                                                                                                                                                              /product= "endostatin'
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                                                                                                                                                                                                                                                                                                                                                                                                            AAC88289 standard; DNA; 552
                                                                                                                                                                          Endostatin; antiangiogenic; angiogenesis; human; inflammation; angiogenesis-dependent disease; ds.
                                                                                                                                                                                                                                                         Human
                                                                                                                                                                                                                                                                                                           02-MAR-2001
                                                                                                                                                                                                                                                                                                                                                          AAC88289;
                                                                                                                                Homo sapiens
                               16-NOV-2000
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Pred. No. 5.1e-102;
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention provides endostatin peptides which can be used in the modulation of angiogenesis. This is useful in the treatment of cancers, inflammation, rheumatoid arthritis, chronic articular rheumatism, psoriasis, disorders associated with inopportune invasion of prematurity, macular degeneration, corneal graft rejection, of prematurity, macular degeneration, corneal graft rejection, retrolental fibroplasia, rubeosis, capillary proliferation in atherosclerotic plaques and osteoporosis. Other angiogenesis-dependent diseases include Osler-Webber syndrome, myocardial angiogenesis, plaque neovasculisation, telangiectasia, haemophilac joints and wound granulation. In addition, the peptides can be used as birth control accerts.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Endostatin peptide comprising at least four endostatin amino acid residues are e.g. angiogenesis inhibitors for treating cancer and diabetic retinopathy.
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AGTGCCGCGAGCTGCCATCACGCCTACATCGTGCTCTGCATTGAGAACAGCTTCATGACT
                                                                    Page 111-112; 146pp; English.
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Pred. No. 5.1e-102;
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                                                                                                                                                                                              regulating angiogenesis related to cancer, arthritis, macular degeneration, and diabetic retinopathy. The compositions are useful for inhibiting angiogenic related diseases. The method and compositions are useful in treating diseases and processes that are mediated by angiogenesis including haemangioma, solid tumours, blood borne tumours, leukaemia, metastasis, telangiectasia, psoriasis, scleroderma, pyogenic granuloma, myocardial angiogenesis (Crohn's disease, plaque neovascularisation, coronary collaterals, cerebral collaterals, atteriovenous malformations, ischaemic limb angiogenesis, corneal diseases, rubeosis, neovascular glaucoma, diabetic retinopathy,
                                                                retrolental fibroplasia, arthritis, diabetic neovascularisation, macular degeneration, wound healing, peptic ulcer, Helicobacter related diseases, fractures, keloids, vasculogenesis, haematopoiesis, ovulation, menstruation, placentation, and cat scratch fever. The method of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention relates to methods and compositions of inhibiting anglogenesis. The method comprises administering to a human or animal a composition comprising a protein that is homologous to PITSLRE protein kinases (PK) and an anglogenic factor. The method is useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Regulating angiogenesis for treating scleroderma, leukemia, keloids ladministering a protein that is homologous to PITSLRE protein kinase an angiogenic factor or a protein kinase and its active fragments.
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/note= "This sequence lacks a start codon"
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invention provides a therapy for cancer that has minimal side effects. The present sequence encodes human endostatin which is used to generat angiogenesis-inhibiting peptides

to generate

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                        (KANG-)
                                            18-JAN-2001; 2001CN-00102221
                                                                                              25-JUL-2001
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The present invention relates to a PCR based method, which can be used to CC obtain the gene of human vascular endothelial inhibin from a human CC obtain the gene of human vascular endothelial inhibin from a human CC mammary gland cDNA library by cloning, and the yeast expression plasmid CC ppICZAA-Endo containing the gene and its transformed pichia engineering CC expression is used to prepare recombined human vascular endothelial CC expression is used to prepare recombined human vascular endothelial CC expression is used to prepare recombined human vascular endothelial collevels, easy large-scale production, low cost and the modification CC mechanism after protein maturation, folding and translation. The human CC vascular endothelial inhibin has the bioactivity to suppress the CC reproduction of vascular endothelial cells and the growth of tumor, so it CC can be used as anticancer medicine. The present sequence is a coding sequence described in the exemplification of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              This invention describes a novel preparation which inhibits the proliferation of blood vessel endothelium and prevents the regeneration activity of tumour blood vessels. The preparation can also be used as a biological preparation in the treatment of tumours. This sequence encode the human endostatin gene described in the invention
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Pred. No. 5.1e-102;
Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               96 T; 0 U; 0 Other;
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RESULT 14
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                                                                                                                                                                             Matches
                                                                                                                                                                                          Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                   A method has been developed for the production of endostatin (I). The method comprises: (i) culturing host cells expressing the gene for (I); (ii) recovering the gene product; (iii) refolding it at near neutral pH; and (iv) isolating properly folded (I). (I) are inhibitors of angiogenesis, by inhibiting migration and proliferation of endothelial cells. The method is used to produce human or murine (I) from inclusion bodies formed in recombinant bacteria. (I), the C-terminal fragment of type alpha 1 collagen XVIII, is known to inhibit growth of some metastases and primary tumours. The present sequence represents a cDNA isolate containing a PCR fragment encoding human endostatin, including the native human collagen XVIII C-terminal region
                                                                                                                                                                                                                                       Sequence 563 BP; 87 A; 198 C; 181 G; 97 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 49; Page 47-48; 52pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Production of endostatin by refolding recombinant protein, useful as anticancer and antimetastatic agent.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Endostatin; inclusion body; expression; anticancer; antimetastatic; angiogenesis; migration; proliferation; inhibition; collagen XVIII;
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                                                                                                                                                                         Conservative
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                                                                                                                                                                 100.0%; Score 546; DB 2;
100.0%; Pred. No. 5.1e-102;
tive 0; Mismatches 0;
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25-FEB-1999;
18-NOV-1999;
                    P-PSDB; AAB08407
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99US-0121633P.
99US-0166176P.
                                                                 Wang L,
                                                                                                                                                                                                                                                                                                                                                                              Location/Qualifiers
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Angiogenesis inhibitor; KED; kringle protein; angiogenesis; tPA; tissue plasminogen activator; tumour; atheroselerosis; arthritis; retinopathy; bronchial vascular congestion; inflammatory bowel disea adult respiratory distress syndrome; Castleman's disease; psoriasis; hepatitis; aneurysm; renal disease; haemangioma; ss.
                                                                                                                                                                                                                                                                                                                                                                                                  DNA encoding a human angiogenesis inhibitor designated KED
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/product= "angiogenesis inhibitor KED"
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Query Match 100.0%;
Best Local Similarity 100.0%;
Matches 546; Conservative 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Use of kringle protein and kringle derived from plasminogen and composition comprising kringle proteins for treating tumor and atherosclerosis, arthritis and retinopathy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 816 BP; 156 A; 259 C; 254 G; 147 T; 0 U; 0 Other;
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805
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Pred. No. 5.2e-102;
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